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(54) Title: AMINE-DERIVATIZED NUCLEOSIDES AND OLIGONUCLEOSIDES

(57) Abstract

Nucleosides and oligonucleosides functionalized to include alkylamino functionality, and derivatives thereof. In certain embodiments, the compounds of the invention further include steroids, reporter molecules, reporter enzymes, lipophilic molecules, peptides or proteins attached to the nucleosides through the alkylamino group.

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AMINE-DERIVATIZED NUCLEOSIDES AND OLIGONUCLEOSIDES

CROSS REFERENCE TO RELATED APPLICATIONS

This application is a continuation-in-part of United States Application Serial No. 08/117,363, filed September 3, 1994 and of Application PCT/US91/00243, filed January 11, 1991 (now United States Application Serial No. 07/854,634, filed July 1, 1992). Application Serial No. 08/117,363, in turn, is a continuation-in-part of Application PCT/US92/09196, filed October 23, 1992, which is a continuation-in-part of United States Application Serial No. 07/782,374, filed October 24, 1991. Application Serial No. 07/782,374 and Application PCT/US91/00243 are both continuations-in-part of United States Application Serial Nos. 07/463,358, filed January 11, 1990, and 07/566,977, filed August 13, 1990. The entire disclosures of each of these applications, which are assigned to the assignee of this application, are incorporated herein by reference.

FIELD OF THE INVENTION

This application is directed to nucleosides, oligonucleotides and oligonucleosides functionalized to include alkylamino functionality, and derivatives thereof. In certain embodiments, the compounds of the invention further include steroids, reporter molecules, reporter enzymes, lipophilic molecules, peptides or proteins attached to the nucleosides through the alkylamino group.

25 BACKGROUND OF THE INVENTION

Messenger RNA (mRNA) directs protein synthesis. Antisense methodology is the complementary hybridization of

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relatively short oligonucleotides to mRNA or DNA such that the normal, essential functions of these intracellular nucleic acids are disrupted. Hybridization is the sequence-specific hydrogen bonding via Watson-Crick base pairs of oligonucleo-5 tides to RNA or single-stranded DNA. Such base pairs are said to be complementary to one another.

naturally occurring events that provide disruption of the nucleic acid function, discussed by Cohen in Oligonucleotides: Antisense Inhibitors of Gene Expression, CRC 10 Press, Inc., Boca Raton, Florida (1989) are thought to be of The first, hybridization arrest, denotes the terminating event in which the oligonucleotide inhibitor binds to the target nucleic acid and thus prevents, by simple steric hindrance, the binding of essential proteins, most often 15 ribosomes, to the nucleic acid. Methyl phosphonate oligonucleotides (Miller, et al., Anti-Cancer Drug Design 1987, 117) and α -anomer oligonucleotides are the two most extensively studied antisense agents which are thought to disrupt nucleic acid function by hybridization arrest.

The second type of terminating event for antisense oligonucleotides involves the enzymatic cleavage of the targeted RNA by intracellular RNase H. A 2'-deoxyribofuranosyl oligonucleotide or oligonucleotide analog hybridizes with the targeted RNA and this duplex activates the RNase H enzyme to cleave the RNA 25 strand, thus destroying the normal function of the RNA. Phosphorothicate oligonuclectides are the most prominent example of an antisense agent that operates by this type of antisense terminating event.

Considerable research is being directed the 30 application of oligonucleotides and oligonucleotide analogs as antisense agents for diagnostics, research reagents and At least for therapeutic potential therapeutic purposes. purposes, the antisense oligonucleotides and oligonucleotide analogs must be transported across cell membranes or taken up 35 by cells to express activity. One method for increasing membrane or cellular transport is by the attachment of a pendant lipophilic group.

Ramirez, et al., J. Am. Chem. Soc. 1982, 104, 5483, introduced the phospholipid group 5'-O-(1,2-di-O-myristoyl-sn-glycero-3-phosphoryl) into the dimer TpT independently at the 3' and 5' positions. Subsequently Shea, et al., Nuc. Acids 5 Res. 1990, 18, 3777, disclosed oligonucleotides having a 1,2-di-O-hexyldecyl-rac-glycerol group linked to a 5'-phosphate on the 5'-terminus of the oligonucleotide. Certain of the Shea, et. al. authors also disclosed these and other compounds in patent application PCT/US90/01002. A further glucosyl phospholipid was disclosed by Guerra, et al., Tetrahedron Letters 1987, 28, 3581.

In other work, a cholesteryl group was attached to the inter-nucleotide linkage between the first and second nucleotides (from the 3' terminus) of an oligonucleotide. This work is disclosed in United States patent 4,958,013 and further by Letsinger, et al., Proc. Natl. Acad. Sci. USA 1989, 86, 6553. The aromatic intercalating agent anthraquinone was attached to the 2' position of a sugar fragment of an oligonucleotide as reported by Yamana, et al., Bioconjugate Chem. 1990, 1, 319. The same researchers placed pyrene-1-methyl at the 2' position of a sugar (Yamana et. al., Tetrahedron Lett. 1991, 32, 6347).

Lemairte, et al., Proc. Natl. Acad. Sci. USA 1986, 84, 648; and Leonetti, et al., Bioconjugate Chem. 1990, 1, 149.

The 3' terminus of the oligonucleotides each include a 3'-terminal ribose sugar moiety. The poly(L-lysine) was linked to the oligonucleotide via periodate oxidation of this terminal ribose followed by reduction and coupling through a N-morpholine ring. Oligonucleotide-poly(L-lysine) conjugates are described in European Patent application 87109348.0. In this instance the lysine residue was coupled to a 5' or 3' phosphate of the 5' or 3' terminal nucleotide of the oligonucleotide. A disulfide linkage has also been utilized at the 3' terminus of an oligonucleotide to link a peptide to the oligonucleotide as is described by Corey, et al., Science 1987, 238, 1401; Zuckermann, et al., J. Am. Chem. Soc. 1988, 110, 1614; and Corey, et al., J. Am. Chem. Soc. 1989, 111, 8524.

Nelson, et al., Nuc. Acids Res. 1989, 17, 7187 describe a linking reagent for attaching biotin to the 3'-terminus of an oligonucleotide. This reagent, N-Fmoc-O-DMT-3-amino-1,2propanediol is now commercially available from Clontech 5 Laboratories (Palo Alto, CA) under the name 3'-Amine on. It is also commercially available under the name 3'-Amino-Modifier reagent from Glen Research Corporation (Sterling, VA). also utilized to link a peptide to was oligonucleotide as reported by Judy, et al., Tetrahedron 10 Letters 1991, 32, 879. A similar commercial reagent (actually. series of such linkers having various polymethylene connectors) for linking to the 5'-terminus of an oligonucleotide is 5'-Amino-Modifier C6. These reagents are available from Glen Research Corporation (Sterling, VA). These 15 compounds or similar ones were utilized by Krieg, et al., Antisense Research and Development 1991, 1, 161 to link fluorescein to the 5'-terminus of an oligonucleotide. compounds of interest have also been linked to the 3'-terminus of an oligonucleotide. Asseline, et al., Proc. Natl. Acad. 20 Sci. USA 1984, 81, 3297 described linking acridine on the 3'terminal phosphate group of an poly (Tp) oligonucleotide via a Haralambidis, et al., Tetrahedron polymethylene linkage. Letters 1987, 28, 5199 report building a peptide on a solid state support and then linking an oligonucleotide to that 25 peptide via the 3' hydroxyl group of the 3' terminal nucleotide of the oligonucleotide. Chollet, Nucleosides & Nucleotides 1990, 9, 957 attached an Aminolink 2 (Applied Biosystems, Foster City, CA) to the 5' terminal phosphate of an oligonucleotide. They then used the bifunctional linking group 30 SMPB (Pierce Chemical Co., Rockford, Il) to link an interleukin protein to the oligonucleotide.

An EDTA iron complex has been linked to the 5 position of a pyrimidine nucleoside as reported by Dreyer, et al., Proc. Natl. Acad. Sci. USA 1985, 82, 968. Fluorescein has been linked to an oligonucleotide in the same manner as reported by Haralambidis, et al., Nucleic Acid Research 1987, 15, 4857 and biotin in the same manner as described in PCT application

PCT/US/02198. Fluorescein, biotin and pyrene were also linked in the same manner as reported by Telser, et al., J. Am. Chem. Soc. 1989, 111, 6966. A commercial reagent, Amino-Modifier-dT, from Glen Research Corporation (Sterling, VA) can be utilized to introduce pyrimidine nucleotides bearing similar linking groups into oligonucleotides.

Cholic acid linked to EDTA for use in radioscintigraphic imaging studies was reported by Betebenner, et.al., Bioconjugate Chem. 1991, 2, 117; however, it is not known to link cholic acid to nucleosides, nucleotides or oligonucleotides.

OBJECTS OF THE INVENTION

It is one object of this invention to provide 15 nucleosides, oligonucleotides and oligonucleosides that include alkylamino chemical functionality.

It is a further object of the invention to provide compounds having improved transfer across cellular membranes.

It is another object to provide compounds that include 20 intercalators, nucleic acid cleaving agents, cell surface phospholipids, and/or diagnostic agents.

It is yet another object to provide improvements in research and diagnostic methods and materials for assaying bodily states in animals, especially disease states.

It is an additional object of this invention to provide therapeutic and research materials having improved transfer and uptake properties for the treatment of diseases through modulation of the activity of DNA or RNA.

30 BRIEF DESCRIPTION OF THE INVENTION

These and other objects are satisfied by the present invention, which provides compounds containing alkylamino chemical functionality. In one aspect, the invention provides nucleosides having base portions and ribofuranosyl sugar portions. Such nucleosides bear at a 2'-0-position, a 3'-0-position, or a 5'-0-position a substituent having formula:

$$-R_A-N(R_{1a})(R_{1b})$$

where:

 R_A is alkyl having from 1 to about 10 carbon atoms or R_A is $(CH_2-CH_2-Q-)_x$;

 R_{1a} and R_{1b} , independently, are H, R_2 , or an amine protecting group, have formula $C(X) - R_2$, $C(X) - R_A - R_2$, $C(X) - Q - R_A - R_2$, $C(X) - Q - R_2$, or, together, complete a cycloalkyl or cycloaryl moiety having two to six carbon atoms and one or two nitrogen atoms;

R₂ includes a steroid molecule, a reporter molecule, a 10 lipophilic molecule, a reporter enzyme, a peptide, a protein, or has formula -Q-(CH₂-Q-)_x-R₃;

X is O or S;

each Q is, independently, is NH, O, or S;

x is 1 to about 200;

15 R_3 is H, R_A , C(O)OH, C(O)OR_A, C(O)R₄, R_A -N₃, R_A -NH₂, or R_A -SH; and

 R_4 is Cl, Br, I, $SO_2^{\dagger}R_5$ or has structure:

-*S (CH₂)_m

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m is 2 to 7; and

R_s is alkyl having 1 to about 10 carbon atoms.

In another aspect, the invention provides oligonucleotides and oligonucleosides comprising a plurality of linked nucleosides, wherein each nucleoside includes a ribofuranosyl sugar portion and a base portion and at least one (preferably more than one) of the nucleosides bears at a 2'-O-position, a 3'-O-position, or a 5'-O-position a substituent having formula of R_a-N(R_{1a})(R_{1b}).

In another aspect the invention provides methods for preparing oligonucleotides and oligonucleosides comprising the steps of contacting nucleosides according to the invention for a time and under reaction conditions effective to form a covalent bond therebetween. In preferred embodiments, at least one of the nucleosides bears a phosphoramidate group at its 2'-O-position or at its 3'-O-position.

In other embodiments, compounds according to the invention are prepared by contacting a nucleoside,

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oligonucleotide or oligonucleoside with derivatizing reagents. For example, a nucleoside, oligonucleotide or oligonucleoside bearing a 2'-hydroxy group, a 3'-hydroxy group, or a 5'-hydroxy group under basic conditions with a compound having formula L₁-5 R_A-N(R_{1a})(R_{1b}) wherein L₁ is a leaving group such as a halogen and at least one of R_{1a} and R_{1b} is an amine protecting group.

The present invention also provides methods for inhibiting the expression of particular genes in the cells of an organism, comprising administering to said organism a compound according to the invention. Also provided are methods for inhibiting transcription and/or replication of particular genes or for inducing degradation of particular regions of double stranded DNA in cells of an organism by administering to said organism a compound of the invention. Further provided are methods for killing cells or virus by contacting said cells or virus with a compound of the invention. The compound can be included in a composition that further includes an inert carrier for the compound.

DETAILED DESCRIPTION OF THE INVENTION

This invention provides nucleosides, oligonucleotides and oligonucleosides containing alkylamino chemical functionality. The nucleoside subunits can be "natural" or "synthetic" moieties. Each nucleoside is formed from a naturally occurring or synthetic base and a naturally occurring or synthetic pentofuranosyl sugar group.

The term "oligonucleotide" refers to a polynucleotide formed from a plurality of linked nucleotide units. The nucleotides units each include a nucleoside unit. In the context of this invention, the term "oligonucleoside" refers to a plurality of nucleoside units that are linked together. In a generic sense, since each nucleotide unit of an oligonucleotide includes a nucleoside therein, the term "oligonucleoside" can be considered to be inclusive of oligonucleotides (i.e., nucleosides linked together via phosphate linking groups). In a further sense, the term "oligonucleoside" also refers to a plurality of nucleosides

that are linked together via linkages other than phosphate linkages. The term "oligonucleoside" thus effectively includes naturally occurring species or synthetic species formed from naturally occurring subunits. For brevity, the term "oligonucleoside" will be used as encompassing both phosphate linked (oligonucleotides) and non-phosphate linked polynucleoside species.

Oligonucleosides according to the invention also can include modified subunits. Representative modifications 10 include modification of a heterocyclic base portion of a nucleoside or a sugar portion of a nucleoside. modifications are disclosed in the following United States Patent Applications: Serial No. 463,358, filed January 11, 1990, entitled Compositions And Methods For Detecting And 15 Modulating RNA Activity; Serial No. 566,977, filed August 13, 1990, entitled Sugar Modified Oligonucleotides That Detect And Modulate Gene Expression; Serial No. 558,663, filed July 27, 1990, entitled Novel Polyamine Conjugated Oligonucleotides; Serial No. 558,806, filed July 27, 1991, entitled Nuclease 20 Resistant Pyrimidine Modified Oligonucleotides That Detect And Modulate Gene Expression and Serial No. PCT/US91/00243, filed January 11, 1991, entitled Compositions and Methods For Detecting And Modulating RNA Activity. Each of these patent applications are assigned to the assignee of this invention. 25 The disclosure of each is incorporated herein by reference.

The term oligonucleoside thus refers to structures that include modified portions, be they modified sugar moieties or modified base moieties, that function similarly to natural bases and natural sugars. Representative modified bases include deaza or aza purines and pyrimidines used in place of natural purine and pyrimidine bases; pyrimidines having substituent groups at the 5 or 6 position; and purines having altered or replacement substituent groups at the 2, 6 or 8 positions. Representative modified sugars include carbocyclic or acyclic sugars, sugars having substituent groups at their 2' position, and sugars having substituents in place of one or more hydrogen atoms of the sugar. Other altered base moieties

and altered sugar moieties are disclosed in United States Patent 3,687,808 and PCT application PCT/US89/02323.

Altered base moieties or altered sugar moieties also include other modifications consistent with the spirit of this invention. Such oligonucleosides are best described as being structurally distinguishable from yet functionally interchangeable with naturally occurring or synthetic wild type oligonucleotides. All such oligonucleosides are comprehended by this invention so long as they function effectively to mimic the structure of a desired RNA or DNA strand.

For use in antisense methodology, the oligonucleosides of the invention preferably comprise from about 10 to about 30 subunits. It is more preferred that such oligonucleosides comprise from about 15 to about 25 subunits. As will be 15 appreciated, a subunit is a base and sugar combination suitably bound to adjacent subunits through, for example, a phosphorouscontaining (e.g., phosphodiester) linkage or some other linking moiety. The nucleosides need not be linked in any particular manner, so long as they are covalently bound. 20 linkages are those between the 3'- and 5'-positions or 2'- and 5'-positions of adjacent nucleosides. Exemplary linking moieties are disclosed in the following references: Beaucage, et al., Tetrahedron 1992, 48, 2223 and references cited therein; and United States Patent Applications: Serial No. 25 703,619, filed May 21, 1991; Serial No. 903,160, filed June 24, 1992; Serial No. 039,979, filed March 20, 1993; Serial No. 039,846, filed March 30, 1993; and Serial No. 040,933, filed March 31, 1993. Each of the foregoing patent applications are assigned to the assignee of this invention. The disclosure of 30 each is incorporated herein by reference.

It is preferred that the RNA or DNA portion which is to be modulated using oligonucleosides of the invention be preselected to comprise that portion of DNA or RNA which codes for the protein whose formation or activity is to be modulated.

35 The targeting portion of the composition to be employed is, thus, selected to be complementary to the preselected portion

of DNA or RNA, that is, to be an antisense oligonucleoside for that portion.

In accordance with one preferred embodiment of this invention, the compounds of the invention hybridize to HIV mRNA encoding the tat protein, or to the TAR region of HIV mRNA. In another preferred embodiment, the compounds mimic the secondary structure of the TAR region of HIV mRNA, and by doing so bind the tat protein. Other preferred compounds are complementary to sequences for herpes, papilloma and other viruses.

10 The nucleosides and oligonucleosides of the invention can be used in diagnostics, therapeutics and as research reagents They can be used in pharmaceutical compositions by including a suitable pharmaceutically acceptable diluent or They further can be used for treating organisms 15 having a disease characterized by the undesired production of The organism should be contacted with a protein. oligonucleotide having a sequence that is capable of specifically hybridizing with a strand of nucleic acid coding for the undesirable protein. Treatments of this type can be 20 practiced on a variety of organisms ranging from unicellular prokaryotic and eukaryotic organisms to multicellular eukaryotic organisms. Any organism that utilizes DNA-RNA transcription or RNA-protein translation as a fundamental part of its hereditary, metabolic or cellular control is susceptible 25 to therapeutic and/or prophylactic treatment in accordance with the invention. Seemingly diverse organisms such as bacteria, yeast, protozoa, algae, all plants and all higher animal forms, including warm-blooded animals, can be treated. Further, since each cell of multicellular eukaryotes can be treated since they 30 include both DNA-RNA transcription and RNA-protein translation as integral parts of their cellular activity. Many of the organelles (e.g., mitochondria and chloroplasts) of eukaryotic cells also include transcription and translation mechanisms. Thus, single cells, cellular populations or organelles can also 35 be included within the definition of organisms that can be treated with therapeutic or diagnostic oligonucleotides. used herein, therapeutics is meant to include the eradication

of a disease state, by killing an organism or by control of erratic or harmful cellular growth or expression.

In one aspect, the present invention is directed to nucleosides and oligonucleosides that bear at least one aminecontaining substituent at a position. Such substituents preferably have formula -R_A-N(R_{1a})(R_{1b}) and are appended at 2'-O-, 3'-O-, and/or 5'-O-positions.

Each R_A according to the invention is an alkyl moiety independently selected to having 1 to about 10 carbon atoms or 10 R_A is a polyether, a polythioether or polyalkylamine. The term "alkyl" is intended to include straight chain and branched hydrocarbons. The preferred length of these hydrocarbons is 1 to about 7 carbon atoms, more preferably 3 or 4 carbon atoms.

 R_{1a} and R_{1b} according to the invention are H, R_2 , an amine 15 protecting group, have formula $C(X) - R_2$, $C(X) - R_A - R_2$, $C(X) - Q - R_A - R_2$, C(X)-Q-R₂, or, together, complete a cycloalkyl or cycloaryl moiety having two to six carbon atoms and one or two nitrogen atoms such as an imidazole moiety. R_{1a} and R_{1b} according to the invention are H, R_A , R_2 , an amine protecting group, or have 20 formula $C(X) - R_2$, $C(X) - R_3 - R_2$, $C(X) - Q - R_3 - R_2$, $C(X) - Q - R_3$. Protecting groups are known per se as chemical functional groups that can be selectively appended to and removed from functionalities, such as amine groups. These groups are present in a chemical compound to render such functionality inert to chemical 25 reaction conditions to which the compound is exposed. See, e.g., Greene and Wuts, Protective Groups in Organic Synthesis, 2d edition, John Wiley & Sons, New York, 1991. Numerous amine protecting groups are known in the art, including, but not limited to: phthalimide (PHTH), trifluoroacetate (triflate), 30 allyloxycarbonyl (Alloc), benzyloxycarbonyl chlorobenzyloxycarbonyl, t-butyloxycarbonyl fluorenylmethoxycarbonyl (Fmoc), and isonicotinyloxycarbonyl (i-Noc) groups. (see, e.g., Veber and Hirschmann, et al., J. Org. Chem. 1977, 42, 3286 and Atherton, et al., The Peptides, 35 Gross and Meienhofer, Eds, Academic Press; New York, 1983; Vol. 9 pp. 1-38).

 R_2 can include a steroid molecule, a reporter molecule, a lipophilic molecule, a reporter enzyme, a peptide, a protein (i.e., a substituent consisting essentially of same), or a molecule having formula -Q-(CH₂CH₂-Q-)_x-R₃. For the purposes of 5 this invention the terms "reporter molecule" and "reporter enzyme" are inclusive of those molecules or enzymes that have physical or chemical properties that allow them to be identified in gels, fluids, whole cellular systems, broken cellular systems and the like utilizing physical properties 10 such as spectroscopy, radioactivity, colorimetric assays, fluorescence, and specific binding. Steroids include those compounds chemical that contain perhydro-1,2a cyclopentanophenanthrene ring system. Proteins and peptides are utilized in their usual sense as polymers of amino acids. 15 Normally peptides comprise such polymers that contain a smaller number of amino acids per unit molecule than do the proteins. Lipophilic molecules include naturally-occurring and synthetic aromatic and non-aromatic moieties such as fatty acids, esters, alcohols and other lipid molecules, substituted aromatic groups 20 such as dinitrophenyl groups, cage structures such as adamantane and buckminsterfullerenes, and aromatic hydrocarbons benzene, perylene, as phenanthrene, anthracene, naphthalene, pyrene, chrysene, and naphthacene.

Particularly useful as steroid molecules are the bile 25 acids including cholic acid, deoxycholic acid and dehydrocholic acid; steroids including cortisone, digoxigenin, testosterone and cholesterol and even cationic steroids such as cortisone having a trimethylaminomethyl hydrazide group attached via a double bond at the 3 position of the cortisone rings. 30 Particularly useful as reporter molecules dinitrophenyl, and fluorescein dyes. Particularly useful as lipophilic molecules are alicyclic hydrocarbons, saturated and unsaturated fatty acids, waxes, terpenes and polyalicyclic hydrocarbons including adamantane and buckminsterfullerenes. 35 Particularly useful as reporter enzymes are alkaline phosphatase and horseradish peroxidase. Particularly useful as

peptides and proteins are sequence-specific peptides and

proteins including phosphodiesterase, peroxidase, phosphatase and nuclease proteins. Such peptides and proteins include SV40 peptide, RNaseA, RNase H and Staphylococcal nuclease. Particularly useful as terpenoids are vitamin A, retinoic acid, retinal and dehydroretinol.

 R_2 also can have formula $-Q-(CH_2CH_2-Q-)_x-R_3$, where Q is O, S, or NH. Subscript x can be 1 to about 200, preferably about 20 to about 150, more preferably about 10 to about 50. Preferably, Q are selected to be O, such that R_2 constitutes a 10 poly(ethyleneglycol) (PEG) group (i.e., $R_3 = H$) or a functionalized derivative thereof (e.g., $R_3 = C(O)Cl$). R_3 can be H, R_A , C(O)OH, $C(O)OR_A$, $C(O)R_4$, R_A-N_3 , R_A-NH_2 or R_A-SH where R_4 is F, Cl, Br, I, SO_2R_5 or a small thio-containing heterocycle having structure:

- S (CH₂) m

where m is 2 to 7. Representative PEG-containing R₂ groups are disclosed by Ouchi, et al., Drug Design and Discovery 1992, 9, 93, Ravasio, et al., J. Org. Chem. 1991, 56, 4329, and Delgardo et. al., Critical Reviews in Therapeutic Drug Carrier Systems 1992, 9, 249.

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Oligonucleosides according to the invention can be assembled in solution or through solid-phase reactions, for example, on a suitable DNA synthesizer utilizing nucleosides according to the invention and/or standard nucleotide precursors. The nucleosides and nucleotide precursors can already bear alkylamino groups or can be later modified to bear such groups.

In the former case, compounds according to the invention are prepared by, for example, reacting nucleosides bearing at least one free 2'-, 3'-, or 5'-hydroxyl group under basic conditions with a compound having formula L_1 -(CH₂)_n-N(R_{1a})(R_{1b}) where L_1 is a leaving group and at least one of R_{1a} and R_{1b} is an amine protecting group. Displacement of the leaving group through nucleophilic attack of an oxygen anion produces the desired amine derivative. Leaving groups according to the invention include but are not limited to halogen, alkyl-

sulfonyl, substituted alkylsulfonyl, arylsulfonyl, substituted arylsulfonyl, hetercyclcosulfonyl or trichloroacetimidate. A more preferred group includes chloro, fluoro, bromo, iodo, p-(2,4-dinitroanilino)benzenesulfonyl, benzenesulfonyl, methylsulfonyl (mesylate), p-methylbenzenesulfonyl (tosylate), p-bromobenzenesulfonyl, trifluoromethylsulfonyl (triflate), trichloroacetimidate, acyloxy, 2,2,2-trifluoroethanesulfonyl, imidazoleysulfonyl, and 2,4,6-trichlorophenyl, with bromo being preferred.

Suitably protected nucleosides can be assembled into an oligonucleosides according to known techniques. See, e.g., Beaucage, et al., Tetrahedron 1992, 48, 2223.

Oligonucleosides according to the invention also can be prepared by assembling an oligonucleoside and appending alkylamino functionality thereto. For example, oligonucleosides having free hydroxyl groups can be assembled according to known techniques and then reacted with a reagent having formula L₁-(CH₂)_n-N(R_{1a})(R_{1b}). As will be recognized, however, greater selectivity can be achieved in terms of placement of alkylamino functionality within an oligonucleoside by introducing such functionality, as discussed above, on selected nucleosides and then using both the selected nucleosides and other nucleosides to construct an oligonucleoside.

Once assembled, an oligonucleoside bearing one or more 25 groups having formula $-R_A - N'(R_{1a})$ (R_{1b}) wherein at least one of R_{1a} and R_{1b} is a protecting group is treated with reagents effective to remove the protecting group. Once deprotected, the oligonucleoside can be contacted with electrophillic moieties such as, for example, succinimidyl esters and other activated 30 carboxylic acids including C(=0)-O-succinimide and C(=0)-Opentafluorophenyl, isothiocyanates, sulfonyl chlorides, halacetamides, phospholipid carbocyclic acid active esters, ophenanthroline-5-iodoacetamide, fluorescein isothiocyanate, 1pyrene butyric acid-N-hydroxy succinimide ester and carboxylic 35 acid derivatives of PNA (carboxylic acid derivatives of peptide Preferred electrophillic moieties include nucleic acids). cholesteryl-3-hemisuccinate-N-hydroxy succinimide

pyrene-1-butyric acid-N-hydroxy succinimide ester and polyethylene glycol-propionic acid-N-hydroxy succimide ester.

Thus, the invention first builds the desired linked nucleoside sequence in the normal manner on the DNA synthesizer. One or more (preferably two or more) of the linked nucleosides are then functionalized or derivatized with the lipophilic steroid, reporter molecule, lipophilic molecule, reporter enzyme, peptide or protein.

Additional objects, advantages, and novel features of this invention will become apparent to those skilled in the art upon examination of the following examples, which are not intended to be limiting. All oligonucleotide sequences are listed in a standard 5' to 3' order from left to right.

EXAMPLE 1

Oligonucleotides Having 2'-Protected-Amine Terminating Linking
Group

A. Preparation of 5'-Dimethoxytrityl-2'-(O-Pentyl-N-phthalimido)-2'-Deoxyadenosine Phosphoramidite.

To introduce a functionalization at the 2' position of 20 nucleotides within desired oligonucleotide sequences, 5'dimethoxytrityl-2'-(O-pentyl-N-phthalimido)-2'-deoxyadenosine phosphoramidite was utilized to provide a linking group attached to the 2' position of nucleotide components of an oligonucleotide. This compound was synthesized generally in 25 accordance with the procedures of patent application serial numbers US91/00243 and 463,358, identified above, starting from adenosine. Briefly, this procedure treats adenosine with NaH in dimethylformamide (DMF) followed by treatment with N-(5bromopentyl) phthalimide. Further treatment with (CH3) 3SiCl, Ph-30 C(O)-Cl and NH4OH yields N6-benzyl protected 2'-pentyl-Nphthalimido functionalized adenosine. Treatment with DIPA and CH₂Cl₂ adds a DMT blocking group at the 5' position. phosphitylation gives the desired phosphoramidite compound. This compound was utilized in the DNA synthesizer as a 0.09M 35 solution in anhydrous CH3CN. Oligonucleotide synthesis was

carried out in either an ABI 390B or an ABI 394 synthesizer employing the standard synthesis cycles with an extended coupling time of 10 minutes during coupling of Compound 2 into the oligonucleotide sequence. Coupling efficiency of greater 5 than 98% was observed.

B. 2'-Protected-Amine Linking Group Containing Phosphodiester Oligonucleotides

The following oligonucleotides having phosphodiester inter-nucleotide linkages were synthesized:

10 Oligomer 9: 5' TA*G 3';

Oligomer 10: 5' CCA' G 3';

Oligomer 11 (SEQ ID NO:1): 5' GGC TGA* CTG CG 3';

Oligomer 12 (SEQ ID NO:2): CTG TCT CCA TCC TCT TCA CT;

Oligomer 13: CTG TCT CCA' TCC TCT TCA' CT

- wherein A* represents a nucleotide functionalized to incorporate a pentyl-N-phthalimido functionality. Oligomers 12 and 13 are antisense compounds to the E2 region of the bovine papilloma virus-1 (BPV-1). Oligomers 12 and 13 have the same sequence as Oligomer 3 in Application Serial Number 782,374,
- 20 except for the 2' modification. The oligonucleotides were synthesized in either a 10 μmol scale or a 3 x 1 μmol scale in the "Trityl-On" mode. Standard deprotection conditions (30% NH₄OH, 55°C, 24 hours) were employed. The oligonucleotides were purified by reverse phase HPLC (Waters Delta-Pak C₄ 15 μm,
- 25 300A, 25x100 mm column equipped with a guard column of the same material). They were detritylated and further purified by size exclusion using a Sephadex G-25 column. NMR analyses by both proton and phosphorus NMR confirmed the expected structure for the Oligomers 9 and 10.

30 C. 2'-Protected-Amine Linking Group Containing Phosphorothioate Oligonucleotides

The following oligonucleotides having phosphorothicate inter-nucleotide linkages were synthesized:

Oligomer 14 (SEQ ID NO:3):

35 $T_sT_sG_s$ $C_sT_sT_s$ $C_sC_sA*_s$ $T_s\overset{1}{C_s}T_s$ $T_sC_sC_s$ $T_sC_sG_s$ $T_sC;$ Oligomer 15 (SEQ ID NO:4):

 $T_sG_sG_s$ $G_sA_sG_s$ $C_sC_sA_s$ $T_sA_sG_s$ $C_sG_sA*_s$ G_sG_sC ; and

- 17 -

Oligomer 16:

 $T_sG_sG_s \ G_sA \star_sG_s \ C_sC_sA \star_s \ T_sA \star_sG_s \ C_sG_sA \star_s \ G_sG_sC$ represents a nucleotide functionalized incorporate a pentyl-N-phthalimido functionality and the 5 subscript "s" represents a phosphorothicate inter-nucleotide backbone linkage. Oligomer 14 is an antisense compound directed to the E2 region of the bovine papilloma virus-1 (BPV-Oligomers 15 and 16 are antisense compounds to ICAM. Oligomer 14 has the same sequence as Oligomer 3 in Application 10 Serial Number 782,374, except for the 2' modification whereas Oligomers 15 and 16 have the same sequence as Oligomer 4 in Application Serial Number 782,374 except for the modification. These oligonucleotides were synthesized as per the method of Example 1(B) except during the synthesis, for 15 oxidation of the phosphite moieties, the Beaucage reagent 3H-1,2-benzodithioate-3-one 1,1-dioxide, Radhakrishnan, et al., J. Am. Chem. Soc. 1990, 112, 1253) was used as a 0.24 M solution in anhydrous CH₃CN solvent. oligonucleotides were synthesized in the "Trityl-On" mode and 20 purified by reverse phase HPLC utilizing the purification procedure of Example 1(B).

D. 2'-O-Methyl Derivatized, 2'-Protected-Amine Linking Group Containing RNA Oligonucleotides

The following oligonucleotides having 2'-O-methyl groups on each nucleotide not functionalized with a 2'-protected amine functionalization were synthesized:

Oligomer 17 (SEQ ID NO:24): CCA A*GC CUC AGA; and

Oligomer 18 (SEQ ID NO:25): CCA GGC UCA GA*T

wherein A* represents a nucleotide functionalized to

30 incorporate a pentyl-N-phthalimido functionality and where the
remaining nucleotides except the 3'-terminus nucleotide are
each 2'-O-methyl derivatized nucleotides. The 3'-terminus
nucleotide in both Oligomers 17 and 18 is a 2'-deoxy
nucleotide. Both Oligomers 17 and 18 are antisense compounds

35 to the HIV-1 TAR region. The oligonucleotides were synthesized
as per the method of Example 6 in Application Serial Number

782,374 (utilizing Compound 2 thereof) for introduction of the

nucleotides containing the pentyl-N-phthalimido functionality and appropriate 2-O-methyl phosphoramidite nucleotides from Chemgenes Inc. (Needham, MA) for the remaining RNA nucleotides. The 3'-terminus terminal 2'-deoxy nucleotides were standard phosphoamidites utilized for the DNA synthesizer. The oligonucleotides were deprotected and purified as per the method of Example 1(B).

EXAMPLE 2

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Functionalization Of Oligonucleotides At the 2' Position

A. Functionalization with Biotin

1. Single Site Modification

About 10 O.D. units (A₂₆₀) of Oligomer 12 (see, Example 1) (approximately 60 nmols based on the calculated extinction coefficient of 1.6756 x 10⁵) was dried in a microfuge tube.

15 The oligonucleotide was dissolved in 200 µl of 0.2 M NaHCO₃ buffer and D-biotin-N-hydroxysuccinimide ester (2.5 mg, 7.3 µmols) (Sigma, St. Louis, MO) was added followed by 40 µl DMF. The solution was let stand overnight. The solution was applied to a Sephadex G-25 column (0.7 x 15 cm) and the oligonucleotide fractions were combined. Analytical HPLC showed nearly 85% conversion to the product. The product was purified by HPLC (Waters 600E with 991 detector, Hamilton PRP-1 column 0.7 x 15 cm; solvent A: 50 mM TEAA pH 7.0; B: 45 mM TEAA with 80% acetonitrile: 1.5 ml flow rate: Gradient: 5% B for first 5 minutes, linear (1%) increase in B every minute thereafter) and further desalted on Sephadex G-25 to give the oligonucleotide:

Oligomer 19 (SEQ ID NO:5): CTG TCT CCA* TCC TCT TCA CT wherein A* represents a nucleotide functionalized to incorporate a biotin functionality linked via a 2'-O-pentyl-30 amino linking group to the 2' position of the designated nucleotide. HPLC retention times are shown in Table 1 below.

2. Multiple Site Modification

About 10 O.D. units (A_{260}) of Oligomer 13 (see, Example 1, approximately 60 nmols) was treated utilizing the method of 35 Example 8(A)(1) in Application Serial Number 782,374 with D-biotin-N-hydroxysuccinimide ester (5 mg) in 300 μ l of 0.2 M

NaHCO₃ buffer/ 50 μ l DMF. Analytical HPLC showed 65% of double labeled oligonucleotide product and 30% of single labeled products (from the two available reactive sites). HPLC and Sephadex G-25 purification gave the oligonucleotide:

oligomer 20: CTG TCT CCA' TCC TCT TCA' CT
wherein A' represents nucleotides functionalized to incorporate
a biotin functionality linked via a 2'-O-pentyl-amino linking
group to the 2' position of the designated nucleotide. HPLC
retention times for this product (and its accompanying singly
labeled products) are shown in Table 1 below.

B. Functionalization with Fluorescein

1. Single Site Modification

A 1M Na₂CO₃/1M NaHCO₃ buffer (pH 9.0) was prepared by adding 1M NaHCO, to 1 M Na₂CO₃. A 200 μ l portion of this buffer 15 was added to 10 O.D. units of Oligomer 12 (see, Example 1) in a microfuge tube. A 10 mg portion of fluorescein-isocyanate in 500 μ l DMF was added to give a 0.05 M solution. portion of the fluorescein solution was added to the oligonucleotide solution in the microfuge tube. 20 covered with aluminum foil and let stand overnight. reaction mixture was applied to a Sephadex G-25 column (0.7 x 20 cm) that had been equilibrated with 25% (v/v) ethyl alcohol The column was eluted with the same solvent. in water. Product migration could be seen as a yellow band well separated 25 from dark yellow band of the excess fluorescein reagent. fractions showing absorption at 260 nm and 485 nm were combined and purified by HPLC as per the purification procedure of Example 2(A)(1). Analytical HPLC indicated 81% of the desired doubly functionalized oligonucleotide. The product 30 lyophilized and desalted on Sephadex to give the oligonucleotide:

Oligomer 21 [(SEQ ID NO:9)]: CTG TCT CCA* TCC TCT TCA CT wherein A* represents a nucleotide functionalized to incorporate a fluorescein functionality linked via a 2'-0-35 pentyl-amino linking group to the 2' position of the designated nucleotide. HPLC retention times are shown in Table 1 below.

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2. Multiple Site Modification

A 10 O.D. unit (A₂₆₀) portion of Oligomer 13 (from Example 1) was dissolved in 300 µl of the 1M Na₂HCO₃/ 1M Na₂CO₂ buffer of Example 2(B)(1) and 200 µl of the fluorescein-isothiocyanate stock solution of Example 2(B)(1) was added. The resulting solution was treated as per Example 2(B)(1). Analytical HPLC indicated 61% of doubly labeled product and 38% of singly labeled products. Work up of the reaction gave the oligonucleotide:

Oligomer 22: CTG TCT CCA* TCC TCT TCA* CT wherein A* represents nucleotides functionalized to incorporate a fluorescein functionality linked via a 2'-O-pentyl-amino linking group to the 2' position of the designated nucleotide. HPLC retention times are shown in Table 1 below.

C. Functionalization with Cholic Acid

1. Single Site Modification

A 10 O.D. unit (A_{260}) portion of Oligomer 12 (see, Example 1) was treated with cholic acid-NHS ester (Compound 1 in Application Serial Number 782,374, 5 mg, 9.9 μ mols) in 200 μ l 20 of 0.2 M NaHCO₃ buffer/ 40 μ l DMF. The reaction mixture was heated for 16 hours at 45°C. The product was isolated as per the method of Example 2(B)(1). Analytical HPLC indicated greater than 85% product formation. Work up of the reaction gave the oligonucleotide:

Oligomer 23: CTG TCT CCA* TCC TCT TCA CT wherein A* represents a nucleotide functionalized to incorporate a cholic acid functionality linked via a 2'-O-pentyl-amino linking group to the 2' position of the designated nucleotide. HPLC retention times are shown in Table 1 below.

2. Multiple Site Modification

A 10 O.D. unit (A₂₆₀) portion of Oligomer 13 (see, Example 1) was treated with cholic acid-NHS ester (Compound 1 in Application Serial Number 782,374, 10 mg, 19.8 μmols) in 300 μl of 0.2 M NaHCO, buffer/ 50 μl DMF. The reaction mixture was heated for 16 hours at 45°C. The product was isolated as per the method of Example 2(A)(1). Analytical HPLC revealed 58% doubly labeled product, 17% of a first singly labeled product

and 24% of a second singly labeled product. Work up as per Example 2(A)(1) gave the oligonucleotide:

Oligomer 24: CTG TCT CCA* TCC TCT TCA* CT
wherein A* represents nucleotides functionalized to incorporate
a cholic acid functionality linked via a 2'-O-pentyl-amino
linking group to the 2' position of the designated nucleotide.
HPLC retention times are shown in Table 1 below.

D. Functionalization with Digoxigenin

1. Single Site Modification

10 A 10 O.D. unit (A_{260}) portion of Oligomer 12 (see, Example 1) was treated with digoxigenin-3-O-methylcarbonyl- ϵ -aminocaproic N-hydroxy succinimide ester (Boehringer Mannheim Corporation, Indianapolis, IN) in 200 μ l of 0.1 M borate pH 8.3 buffer/ 40 μ l DMF. The reaction mixture was let stand 15 overnight. The product was isolated as per the method of Example 2(A)(1). Work up of the reaction gave the oligonucleotide:

Oligomer 25: CTG TCT CCA* TCC TCT TCA CT
wherein A* represents a nucleotide functionalized to
10 incorporate a digoxigenin functionality linked via a 2'-Opentyl-amino linking group to the 2' position of the designated
nucleotide. HPLC retention times are shown in Table 1 below.

2. Multiple Site Modification

A 10 O.D. units (A_{260}) portion of Oligomer 13 (see, 25 Example 1) was treated with digoxigenin-3-0-methylcarbonyl- ϵ -aminocaproic N-hydroxy succinimide ester (Boehringer Mannheim Corporation, Indianapolis, IN) in 300 μ l of 0.1 M borate pH 8.3 buffer/50 μ l DMF. The reaction mixture was let stand overnight. The product was isolated as per the method of 30 Example 2(A)(1). Work up as per Example 2(A)(1) gave the oligonucleotide:

Oligomer 26: CTG TCT CCA* TCC TCT TCA* CT
wherein A* represents nucleotides functionalized to incorporate
a cholic acid functionality linked via a 2'-O-pentyl-amino
linking group to the 2' position of the designated nucleotide.
HPLC retention times are shown in Table 1 below.

TABLE 1

HPLC Retention Times Of Oligonucleotides Functionalized

At 2' Position

5	Oligomer		ntion Time titution M		Substitution	
	Oligomer 12 ¹	21.7	8			
	Oligomer 131		1	22.50		
	Oligomer 192		8			
10	Oligomer 202		1	24.16ª	25.19 ^b	
	Oligomer 213	26.6	5			
	Oligomer 223			26.99ª	29.33 ^b	
	0119001			27.55ª		•
	Oligomer 234	30.1	0 .			
15	Oligomer 24			30.38ª	37.00 ^b	
10	Origomer 21		1	32.22ª	•	
	Oligomer 25	28.0	1			
	Oligomer 26			28.14ª	33.32 ^b	
	Oligomer 20			29.24ª		
20						

- Mono conjugated minor product;
- b Doubly conjugated major product;
- 1 Parent Oligonucleotide no 2' functionalization;
- ² 2' Biotin functionalization;
- 25 3 2' Fluorescein functionalization;
 - 4 2' Cholic Acid functionalization; and
 - ⁵ 2' Digoxigenin functionalization.

The conditions employed were as follows: Waters 600E with 991 detector, Hamilton PRP-1 column 0.7 x 15 cm; solvent 30 A: 50 mM TEAA pH 7.0; B: 45 mM TEAA with 80% acetonitrile: 1.5 ml flow rate: Gradient: 5% B for first 5 minutes, linear (1%) increase in B every minute thereafter.

PROCEDURE A

Confirmation of Structure of Functionalized Oligonucleotides 35 Containing A Tethered 2'-Amino Moiety

Oligonucleotides of the invention were digested with snake venom phosphodiesterase and calf-intestine alkaline phosphatase to their individual nucleosides. After digestion, the nucleoside composition was analyzed by HPLC. The HPLC 40 analysis established that functionalized nucleotide compounds having the tethered 2'-amino moiety thereon were correctly incorporated into the oligonucleotide.

Snake venom phosphodiesterase [Boehringer-Mannheim cat. #108260, 1 mg (1.5 units)/0.5 ml] and alkaline phosphatase from calf intestine (1 unit/microliter, Boehringer-Mannheim cat. # 713023) in Tris-HCl buffer (pH 7.2, 50 mM) were used 5 to digest the oligonucleotides to their component nucleosides. To 0.5 O.D. units of oligonucleotide in 50 μ l buffer (nearly 40 μM final concentration for a 20 mer) was added 5 μl of snake venom phosphodiesterase (nearly 0.3 units/mL, final concentration) and 10 μ l of alkaline phosphatase (app. 150 10 units/mL, final concentration). The reaction mixture was Following incubation, the incubated at 37°C for 3 hours. reaction mixture was analyzed by HPLC using a reverse phase analytical column (app. 30 x 2.5 cm); solvent A: 50 mM TEAA pH 7; solvent B: acetonitrile; gradient 100% for 10 minutes, 15 then 5% B for 15 minutes, then 10% B and then wash. The results of these digestion are shown in Table 2 for representative oligonucleotides.

TABLE 2
OLIGONUCLEOTIDE ANALYSIS VIA ENZYMATIC DIGESTION

20		Observed Ratios**					
	Oligomer	Abs.				267 A*	260 A
	Oligomer 10		2	1		1	
25	Oligomer 11	ί.	3	5	2	1	
	Oligomer 12	I	9	1	8	1	1
	Oligomer 13	! !	9	1	8	2	

Nucleoside having 2'-O-linker attached thereto; and Corrected to whole numbers.

As is evident from Table 2, the correct nucleoside ratios are observed for the component nucleotides of the test oligonucleotides.

PROCEDURE B

Determination of Melting Temperatures (Tm's) of Cholic Acid Oligonucleotide Conjugates

The relative ability of oligonucleotides to bind to 5 their complementary strand is compared by determining the melting temperature of the hybridization complex of the oligonucleotide and its complementary strand. The melting temperature (Tm), a characteristic physical property of double helices, denotes the temperature in degrees centigrade at 10 which 50% helical versus coil (un-hybridized) forms are present. Tm is measured by using the UV spectrum to determine the formation and breakdown (melting) of hybridization. Base stacking, which occurs during hybridization, is accompanied absorption (hypochromicity). ŪΫ reduction in by 15 Consequently a reduction in UV absorption indicates a higher The higher the Tm, the greater the strength of the binding of the strands. Non-Watson-Crick base pairing has a strong destabilizing effect on the Tm. Consequently, absolute fidelity of base pairing is necessary to have optimal binding 20 of an antisense oligonucleotide to its targeted RNA.

1. Terminal End Conjugates

a. Synthesis

A series of oligonucleotides were synthesized utilizing un-functionalized (for synthetic procedures standard 25 oligonucleotides) or the procedure of Example 3(A) Application Serial Number 782,374 for oligonucleotides having a 5'-terminus amino linker bearing oligonucleotide or the procedure of Example 3 (B) in Application Serial Number 782,374 for 5'-terminus cholic acid-bearing oligonucleotides. Each 30 of the oligonucleotides had the following 5-LO antisense sequence: 5' TCC AGG TGT CCG CAT C 3'(SEQ ID NO:6). nucleotides were synthesized on a 1.0 μ mol scale. Oligomer 32 was the parent compound having normal phosphodiester inter-Oligomer 33 incorporated nucleotide linkages. 35 phosphorothicate inter-nucleotide linkages in the basic oligonucleotide sequence. Oligomer 34 is a an intermediate oligonucleotide having a 5'-aminolink at the 5'-terminus of

the basic oligonucleotide sequence and Oligomer 35 was a similar 5'-aminolink compound incorporating phosphorothicate inter-nucleotide linkages. Oligomer 36 is a 5'-terminus cholic acid conjugate of the basic phosphodiester oligonucleo-5 tide sequence while Oligomer 37 is a similar 5'-cholic acid conjugate incorporating phosphorothicate inter-nucleotide linkages. Oligomers 32 and 33 were synthesized in a "Trityl-On" mode and were purified by HPLC. Oligomers 34 and 35 were synthesized as per Example 30(A) in Application Serial Number 10 782,374 without or with Beaucage reagent treatment, to yield phosphodiester or phosphorothicate inter-nucleotide linkages, respectively. Oligomers 36 and 37 were prepared from samples of Oligomers 34 and 35, respectively, utilizing a solution of cholic acid N-hydroxysuccinimide ester (Compound 1) 15 dissolved in DMF as per Example 3(B) in Application Serial Number 782,374. Oligomers 36 and 37 were purified by HPLC. The products were concentrated and desalted in a Sephadex G-25 column. Gel electrophoresis analyses also confirmed a pure product with the pure conjugate moving slower than the parent 20 oligonucleotide or 5'-amino functionalized oligonucleotide.

b. Melting Analysis

The test oligonucleotides [either the phosphodiester, phosphorothicate, cholic acid conjugated phosphodiester, cholic acid conjugated phosphorothicate or 5'-aminolink phosphodiester or phosphorothioate 25 intermediate oligonucleotides of the invention or otherwise] and either the complementary DNA or RNA oligonucleotides were incubated at a standard concentration of 4 μM for each oligonucleotide in buffer (100 mM NaCl, 10 mM Na-phosphate, pH 7.0, 0.1 mM EDTA). 30 Samples were heated to 90 degrees C and the initial absorbance a Guilford Response spectrophotometer II (Corning). Samples were then slowly cooled to 15 degrees C and then the change in absorbance at 260 nm was monitored during the heat denaturation procedure. The temperature was 35 elevated 1 degree/absorbance reading and the denaturation profile analyzed by taking the 1st derivative of the melting Data was also analyzed using a two-state linear curve.

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regression analysis to determine the Tm's. The results of these tests are shown in Table 3 as are the HPLC retention times of certain of the test compounds.

TABLE 3

Melting Temperature Of The Hybridization Complex Of The Oligonucleotide And Its Complementary Strand

Oligomer		Tm **		HPLC Ret. Time*		
		DNA	RNA	minutes		
10 -	32	62.6	62.0			
	33	55.4	54.9			
	34	ND	ND	13.6		
	35	ND	ND	17.0		
15	36	63.4	62.4	22.0		
	37	56.3	55.8	22.5		

- * HPLC conditions: Walters Delta Pak C-18 RP 2.5u column; at 0 min 100% 0.1 TEAA; at 30 min 50% TEAA and 50% Acetonitrile: Flow rate 1.0 ml/min.
 - ** Tm at $4\mu\text{M}$ each strand from fit of duplicate melting curves to 2-state model with linear sloping base line. Conditions: 100 mM NaCl, 10 mM Phosphate, 0.1 mM EDTA, pH 7.0.

25 ND = not determined

As is evident from Table 2, conjugates of cholic acid at the end of the oligonucleotide do not affect the Tm of the oligonucleotides.

2. Strands Incorporating 2'-O-Pentylamino Linker

a. Synthesis

An oligonucleotide of the sequence:

Oligomer 38 (SEQ ID NO:7): GGA* CCG GA*A* GGT A*CG A*G wherein A* represents a nucleotide functionalized to incorporate a pentylamino functionality at its 2'-position was synthesized in a one micromole scale utilizing the method of Example 1(B). The oligonucleotide was purified by reverse phase HPLC, detritylated and desalted on Sephadex G-25. PAGE gel analysis showed a single band. A further oligonucleotide, Oligomer 39, having the same sequence but without any 2'-O-40 amino linker was synthesis in a standard manner. A complementary DNA oligonucleotide of the sequence:

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Oligomer 40 (SEQ ID NO:8): CCT GGC CTT CCA TGC TC was also synthesized in a standard manner as was a complementary RNA oligonucleotide of the sequence:

Oligomer 41 (SEQ ID NO:9): CCU GGC CUU CCA UGC UC.

b. Melting Analysis

Melting analysis was conducted as per the method of Procedure B(1)(b). The results are shown in Table 4.

TABLE 4

Melting Temperature Of The Hybridization Complex Of
The Oligonucleotide And Its Complementary Strand

Oligomer		` T m⁺	
	0119001	DNA ¹ RNA ²	
	38	54.5 58.0	
15	39	60.6 56.9	

Tm at $4\mu\text{M}$ each strand from fit of duplicate melting curves to 2-state model with linear sloping base line. Conditions: 100 mM NaCl, 10 mM Phosphate, 0.1 mM EDTA, pH 7.0.

Against DNA complementary strand, Oligomer 40.

2 Against RNA complementary strand, Oligomer 41

As is evident from Table 4 against the RNA complementary strand the change in Tm's between the strand 25 having 2'-amino linkers thereon and the unmodified strand is 1.1 degrees (0.22 change per modification). Against the DNA strand, the change is -6.1 degrees (-1.2 change per modification). When compared to the parent unmodified oligonucleotide the 2'-amino linker- containing strand has a stabilizing effect upon hybridization with RNA and a destabilizing effect upon hybridization with DNA.

Compounds of the invention were tested for their ability to increase cellular uptake. This was determined by judging either their ability to inhibit the expression of 35 bovine papilloma virus-1 (BPV-1) or an assay involving luciferase production (for HIV-1).

PROCEDURE C

Determination of Cellular Uptake Judged By The Inhibition Of Expression of Bovine Papilloma Virus-1 (bpv-1) As Measured By an E2 Transactivation Assay

For this test, a phosphorothicate oligonucleotide analog of the sequence:

Oligomer 42: CTG TCT CCA TCC TCT TCA CT was used as the basic sequence. This sequence is designed to be complementary to the translation initiation region of the 10 E2 gene of bovine papilloma virus type 1 (BPV-1). Oligomer 42 served as the positive control and standard for the assay. Oligomer 3 (from Example 4 in Application Serial Number 782,374) served as a second test compound. It has the same basic sequence except it is a phosphorothicate oligonucleotide 15 and further it has a cholic acid moiety conjugated at the 3'end of the oligonucleotide. Oligomer 2 (from Example 2 in Application Serial Number 782,374) served as a third test Again it is of the same sequence, it is a compound. phosphorothicate oligonucleotide and it has a cholic acid 20 moiety conjugated at the 5'-end. Oligomer 5 (from Example 5 in Application Serial Number 782,374) served as a fourth test Once again it has the same sequence, is a compound. phosphorothicate oligonucleotide and it has a cholic acid moiety conjugated at both the 3'-end and 5'-end. A fifth test 25 compound was a phosphorothicate oligonucleotide with no significant sequence homology with BPV-1. A sixth test compound was a further phosphorothicate oligonucleotide with no significant sequence homology with BPV-1. The last test compound, the seventh test compound, was a phosphorothicate 30 oligonucleotide with cholic acid conjugated to the 3'-end but having no significant sequence homology with BPV-1. Compounds five, six and seven served as negative controls for the assay.

For each test I-38 cells were plated at 5x10⁴ cells per cm² in 60 mm petri dishes. Eight hours after plating, medium 35 was aspirated and replaced with medium containing the test oligonucleotide and incubated overnight. Following incubation, medium was aspirated and replaced with fresh

medium without oligonucleotide and incubated for one hour. Cells were then transfected by the CaPO₄ method with 2 μg of pE2RE-1-CAT. After a four hour incubation period cells were glycerol shocked (15% glycerol) for 1 minute followed by 5 washing 2 times with PBS. Medium was replaced with DMEM containing oligonucleotide at the original concentration. Cells were incubated for 48 hours and harvested. Cell lysates were analyzed for chloramphenicol acetyl transferase by standard procedures. Acetylated and nonacetylated ¹⁴C-10 chloramphenicol were separated by thin layer chromatography and quantitated by liquid scintillation. The results are expressed as percent acetylation.

Two lots of the positive control compound were found to acetylate at a level of 29% and 30%. The negative 15 controls, test compounds five, six and seven, were found to acetylate at 59%, 58% and 47%, respectively. The 3'-cholic acid conjugate test compound, Oligomer 3, was found to acetylate to 23%, the 5'-cholic acid conjugate test compound, Oligomer 2, was found to acetylate to 36% and the test 20 compound conjugated at both the 3'-end and the 5'-end, Oligomer 5, was found to acetylate to 27%.

The results of this test suggests that placement of a cholic acid moiety at the 3'-terminus of an oligonucleotide increase the activity. This in turn suggests that the 25 increased activity was the result of increased cellular membrane transport.

PROCEDURE D

Determination of Cellular Uptake Judged By Inhibition of pHIVluc With Cholic Acid Linked 2'-O-Methyl Substituted 30 Oligonucleotides

For this test the absence of an oligonucleotide in a test well served as the control. All oligonucleotides were tested as 2'-O-methyl analogs. For this test an oligonucleotide of the sequence:

where each of the nucleotides of the oligonucleotide includes a 2'-O-methyl substituent group served as the basic test compound. The second test compound of the sequence:

Oligomer 44: 5'-CHA CCC AGG CUC AGA

5 wherein CHA represents cholic acid and where each of the nucleotides of the oligonucleotide includes a 2'-0-methyl substituent group, was also of the same sequence as the first test compound. This second test compound included cholic acid conjugated to its 5'-end and was prepared as per the method 10 of Example 3 in Application Serial Number 782,374 utilizing 2'-0-methyl phosphoramidite intermediates as identified in Example 1(C). The third test compound of the sequence:

Oligomer 45: CCC AGG CUC AGA 3'-CHA

wherein CHA represents cholic acid and where each of the nucleotides of the oligonucleotide includes a 2'-O-methyl substituent group was also of the same sequence as the first test compound. The third test compound included cholic acid conjugated to its 3'-end and was prepared as per the method of Example 4 in Application Serial Number 782,374 utilizing 2'-O-methyl phosphoramidite intermediates as identified in Example 1(C). The fourth test compound was a 2'-O-Me oligonucleotide of a second sequence:

Oligomer 46 (SEQ ID NO:11): GAG CUC CCA GGC
where each of the nucleotides of the oligonucleotide includes
25 a 2'-O-methyl substituent group. The fifth test compound was
of sequence:

Oligomer 47: 5'-CHA GAG CUC CCA GGC

wherein CHA represents cholic acid and where each of the nucleotides of the oligonucleotide includes a 2'-O-methyl 30 substituent group. It was of the same sequence as the fifth test compound. This test compound included cholic acid conjugated to its 5'-end and was prepared as per the method of Example 3 in Application Serial Number 782,374 utilizing 2'-O-methyl phosphoramidite intermediates as identified in 35 Example 1(C).

A sixth test compound was a randomized oligonucleotide of the sequence:

Oligomer 48 (SEQ ID NO:12): CAU GCU GCA GCC.

HeLa cells were seeded at 4×10^5 cells per well in 6-Test oligonucleotides were added to well culture dishes. triplicate wells at 1 μM and allowed to incubate at 37°C for 5 20 hours. Medium and oligonucleotide were then removed, cells washed with PBS and the cells were CaPO4 transfected. Briefly, $5\mu g$ of pHIVluc, a plasmid expressing the luciferase cDNA under the transcriptional control of the HIV LTR constructed by ligating the KpnI/HindIII restriction fragments 10 of the plasmids pT3/T7luc and pHIVpap (NAR 19(12)) containing the luciferase cDNA and the HIV LTR respectively, and 6 μg of pcDEBtat, a plasmid expressing the HIV tat protein under the control of the SV40 promoter, were added to 500 μ l of 250 mM $CaCl_2$, then 500 μl of 2x HBS was added followed by vortexing. 15 After 30 minutes, the CaPO4 precipitate was divided evenly between the six wells of the plate, which was then incubated for 4 hours. The media and precipitate were then removed, the cells washed with PBS, and fresh oligonucleotide and media were added. Incubation was continued overnight. Luciferase 20 activity was determined for each well the following morning. Media was removed, then the cells washed 2X with PBS. cells were then lysed on the plate with 200 μl of LB (1% Trit X-100, 25 mM Glycylglycine pH 7.8, 15 mM MgSO4, 4 mM EGTA, 1mM DTT). A 75 μ l aliquot from each well was then added to a well 25 of a 96 well plate along with 75 μ l of assay buffer (25 mM Glycylglycine pH 7.8, 15 mM MgSO4, 4 mM EGTA, 15 mM KPO4, 1 mM DTT, 2.5 mM ATP). The plate was then read in a Dynatec multiwell luminometer that injected 75 μ l of Luciferin buffer (25 mM Glycylglycine pH 7.8, 15 mM MgSO4, 4 mM EGTA, 4 mM DTT, 30 1 mM luciferin) into each well, immediately reading the light emitted (light units).

The random sequence compound (Oligomer 48) and the other non-cholic acid-conjugated test compounds (Oligomers 43 and 46) had comparable activity. The 5'-conjugate of the 35 first sequence (Oligomer 44) also had activity comparable to the non-conjugated compounds. The 5'-conjugate of the second sequence (Oligomer 47) showed a three-fold increase in

activity compared to the non-conjugated compounds and the 3'-conjugate of the first sequence (Oligomer 45) showed a further 3-fold increase in activity compared to Oligomer 47.

All the test cholic acid-bearing oligonucleotides 5 showed significant inhibition of luciferase production compared to non-cholic acid-bearing oligonucleotides. This suggests that the increased activity was the result of increased cellular membrane transport of the cholic acid-bearing test oligonucleotides.

10 EXAMPLE 3

Preparation of 5'-0-[Dimethoxytrityl]-2'-0-[hexyl-(Ω -N-phthalimido)amino]uridine and 5'-0-[dimethoxytrityl]-3'-0-[hexyl(Ω -N-phthalimidoamino)uridine.

2',3'-O-Dibutyl stannylene-uridine was synthesized 15 according to the procedure of Wagner et. al., J. Org. Chem., 1974, 39, 24. This compound was dried over P2O5 under vacuum for 12 hours. To a solution of this compound (29 g, 42.1 mmols) in 200 ml of anhydrous DMF were added (16.8 g, 55 mmols) of 6-bromohexyl phthalimide and 4.5 g of sodium iodide 20 and the mixture was heated at 130°C for 16 hours under argon. The reaction mixture was evaporated, co-evaporated once with toluene and the gummy tar residue was applied on a silica column (500 g). The column was washed with 2L of ethyl acetate (EtOAc) followed by eluting with 10% methanol 25 (MeOH):90% EtOAc. The product, 2'- and 3'-isomers of O-hexyl- Ω -N-phthalimido uridine, eluted as an inseparable mixture $(R_f=0.64 \text{ in } 10\% \text{ MeOH in EtOAc})$. By ^{13}C NMR, the isomeric ration was about 55% of the 2' isomer and about 45% of the 3' isomer. The combined yield was 9.2 g (46.2%). This mixture 30 was dried under vacuum and re-evaporated twice with pyridine. It was dissolved in 150 mL anhydrous pyridine and treated with 7.5 g of dimethyocytrityl chloride (22.13 mmols) and 500 mg of dimethylaminopyridine (DMAP). After 2 hour, thin layer chromatography (TLC; 6:4 EtOAc: Hexane) indicated complete 35 disappearance of the starting material and a good separation between 2' and 3' isomers ($R_f=0.29$ for the 2' isomer and 0.12 for the 3' isomer). The reaction mixture was quenched by the addition of 5mL of CH₃OH and evaporated under reduced pressure. The residue was dissolved in 300mL CH₂Cl₂, washed successively with saturated NaHCO₃ followed by saturated NaCl solution. It was dried over Mg₂SO₄ and evaporated to give 15 g of a brown foam which was purified on a silica gel (500 g) to give 6.5 g of the 2'-isomer and 3.5 g of the 3' isomer.

EXAMPLE 4

Preparation of 5'-0-Dimethoxytrityl-2'-0-[hexyl-(Ω-N-10 phthalimido)amino]uridine-3'-0-(2-cyanoethyl-N,N-diisopropyl)-phosphoramidite.

The 5'-dimethoxytrityl-2'-[O-hexyl-(Ω-N-phthalimido)-amino]uridine (4 g, 5.2 mmole) was dissolved in 40 mL of anhydrous CH₂Cl₂. To this solution diisopropylaminetetra-15 zolide (0.5 g, 2.9 mmol) was added and stirred overnight. TLC (1:1 EtoAC/hexane) showed complete disappearance of starting material. The reaction mixture was transferred with CH₂Cl₂ and washed with saturated NaHCO₃ (100 mL) followed by saturated NaCl solution. The organic layer was dried over 20 anhydrous Na₂SO₄ and evaporated to yield 6.4 g of a crude product which was purified in a silica column (200 g) using 1:1 hexane/EtOAc to give 4.6 g (4.7 mmol, 90%) of the desired phosphoramidite.

EXAMPLE 5

25 Preparation of 5'-0-(Dimethoxytrity1)-3'-0-[hexyl-(Ω-N-phthalimido)amino]uridine-2'-0-succinyl-aminopropyl controlled pore glass.

Succinylated and capped aminopropyl controlled pore glass (CPG; 500Å pore diameter, aminopropyl CPG, 1.0 grams 30 prepared according to Damha et. al., Nucl. Acids Res. 1990, 18, 3813.) was added to 12 ml anhydrous pyridine in a 100 ml round-bottom flask. 1-(3-Dimethylaminopropyl)-3-ethyl-carbodimide (DEC; 0.38 grams, 2.0 mmol)], triethylamine (TEA; 100 µl, distilled over CaH₂), dimethylaminopyridine (DMAP; 0.012 35 grams, 0.1 mmol) and nucleoside 5'-O-dimethoxytrityl-3'-O-

[hexyl-(Ω -N-phthalimidoamino)]uridine (0.6 grams, 0.77 mmol) were added under argon and the mixture shaken mechanically for More nucleoside (0.20 grams) was added and the mixture shaken an additional 24 hours. CPG was filtered off 5 and washed successively with dichloromethane, triethylamine, The CPG was then dried under vacuum, and dichloromethane. suspended in 10 ml piperidine and shaken 15 minutes. was filtered off, washed thoroughly with dichloromethane and again dried under vacuum. |The extent of loading (determined 10 by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm) was approximately 28 μ mol/g. The 5'-O-(dimethoxytrityl)-3'-O-[hexyl-(Ω -N-phthalimidoamino]uridine-2'-0-succinyl-aminopropyl controlled pore glass was used to synthesize the oligomers 5'-GACU*-3' and 5'-15 GCC TTT CGC GAC CCA ACA CU*-3' (SEQ ID NO:13) (where the * indicated the derivatized nucleotide) in an ABI 380B DNA using phosphoramidite chemistry synthesizer 45 and 200 O.D.'s of the 4-mer and 20-mer, conditions. respectively, were obtained from two and three 1 μ mol 20 syntheses after purification by RP-HPLC and desalting.

The oligomer 5'-GACU'-3' was used to confirm the structure of 3'-O-hexylamine tether introduced into the oligonucleotide by NMR. As expected a multiplet signal was observed between 1.0-1.8 ppm in 'H NMR. The oligomer 5'-GCC TTT CGC GAC CCA ACA CU'- 3' belongs to a HCV sequence and it was used to show the nuclease resistance properties of the 3'-O-amino tether [see example 39].

EXAMPLE 6

Preparation of 5'-O-(Dimethoxytrity1)-2'-O-[hexy1-(Ω-N-30 phthalimido)amino] 3'-O-succinyl-aminopropyl controlled pore glass..

The procedure of Example 5 was repeated, except that $5'-0-(Dimethoxytrity1)-2'-0-[hexyl-(<math>\Omega$ -N-phthalimidoamido)-amino]uridine was used in the loading process.

Preparation of 5'-0-(Dimethoxytrityl)-2'-0-(hexylamino)-uridine.

5'-O-(dimethoxytrityl)-2'-O-[hexyl-(Ω-N-phthalimido 5 amino)]uridine (4.5 grams, 5.8 mmol) was dissolved in 200 ml methanol in a 500 ml flask. Hydrazine (1 ml, 31 mmol) was added to the stirring reaction mixture. The mixture was heated to 60-65° in an oil bath and refluxed 14 hours. Solvent was evaporated in vacuo. The residue was dissolved 10 in dichloromethane (250 ml) and extracted twice with an equal volume NH₄OH. The organic layer was evaporated to yield 4.36 grams of crude product, and NMR indicated that the product was not completely pure. R_f=0 in 100% ethyl acetate. The product was used in subsequent reactions without further purification.

15 EXAMPLE 8

Preparation of 5'-0-(dimethoxytrityl)-3'-0-[hexylamino] uridine.

The procedure of Example 7 was repeated, except that $5'-0-(dimethoxytrity1)-3'-0-[hexyl-(<math>\Omega$ -N-phthalimido-amino)] 20 uridine was used as the starting material.

EXAMPLE 9

Preparation of 5'-0- (dimethoxytrityl) -2'-0- [hexyl-N-(1-pyrene propyl carbonyl) amino] uridine

5'-O-Dimethoxytrityl-2'-O-(hexylamino)uridine (0.5g, 25 0.78 mmol) was dissolved in anhydrous DMF (15 mL). 1-Hydroxybenzotriazole (0.16 grams, 1.17 mmol) and 1-pyrene-butyric acid pentafluorophenyl ester (0.53 grams, 1.17 mmol) were added to the reaction mixture. The mixture was stirred under argon at room temperature for 2 hours, after which it 30 was concentrated in vacuo. Residual DMF was coevaporated with toluene. The residue was dissolved in dichloromethane (50 mL) and washed with an equal volume saturated NaHCO3. The aqueous layer was washed with an equal volume saturated NaCl. The 35 aqueous layer was washed with dichloromethane and the combined

organic layers dried over $MgSO_4$ and concentrated. The residue was chromatographed on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product (0.83 grams, 58%) eluted with 100% ethyl acetate (R_f 0.46 by thin-layer chromatography (TLC)).

EXAMPLE 10

Preparation of 5'-O-[Dimethoxytrity1]-2'-O-[hexyl-N-(1-pyrene propyl carbonyl) amino] uridine-3'-O-(2-cyanoethyl-N, N-10 disopropyl) phosphoramidite

5'-0-[Dimethoxytrityl]-2'-0-[hexyl-N-(1-pyrene propyl carbonyl)amino] uridine (0.80 grams, 0.87 mmol) was dissolved in 20 mL dry dichloromethane. 2-Cyanoethyl N, N, N', N'-tetraisopropylphosphorodiamidite (purchased from Sigma Chemical Co; 15 800 μ L, 2.4 mmol) and diisopropylamine tetrazolide (0.090 grams, 0.52 mmol) were added to the mixture, which was stirred under argon for 20 hours | The reaction mixture was then dissolved the residue concentrated in vacuo and The solution was washed with an dichloromethane (75 mL). 20 equal volume of saturated NaHCO3. The aqueous layer was washed with dichloromethane (20 mL) and the combined organic layers washed with an equal volume of saturated NaCl. aqueous layer was washed with dichloromethane (20 mL) and the combined organic layers dried over MgSO4 and concentrated. 25 The residue was chromatographed on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product (750 mg, 78% yield, R_f 0.54 by TLC in 100% ethyl acetate) eluted with 100% ethyl acetate.

30 EXAMPLE 11

Preparation of 2'-0-[hexyl-N-(1-pyrene-propyl-carbonyl) amino] uridine.

5'-O-dimethoxytrityl 2'-O-[hexyl-N-(1-pyrene-propyl-carbonyl)amino]uridine (1.0 g) was dissolved in 20 mL CH₂Cl₂ 35 and kept in ice-bath for 10 minutes. To the cold solution,

5 mL of 80% acetic acid in water was added and the solution was left to stand for 30 minutes. It was then evaporated to dryness and loaded into a silica column and eluted with 10% methanol in methylene chloride to give 2'-O-[hexyl-N-(1-5 pyrene-propyl-carbonyl)amino]uridine.

EXAMPLE 12

Preparation of 5'-0-(dimethoxytrity1)-2'-0-[hexyl-N-(1-pyrene propylcarbonyl)amino]uridine-3'-0-[succinylaminopropyl]-controlled pore glass.

Succinylated/capped aminopropyl controlled pore glass 10 was dried under vacuum for 3 hours immediately before use. A portion (0.3 g) was added to 3 ml anhydrous pyridine in a 50 ml round-bottom flask. DEC (0.12 grams, 0.63 mmol), TEA (25 μ l, distilled over CaH₂), DMAP (0.005 grams, mmol) and 5'-15 O-(dimethoxytrityl)-3'-O-[hexyl-N-(1-pyrene propyl carbonyl] amino]uridine (0.21 grams, 0.22 mmol) were added under argon and the mixture shaken mechanically for 19 hours. nucleoside (0.025 grams) was added and the mixture shaken an additional 5.5 hours. Pentachlorophenol (0.045 grams, mmol) 20 was added and the mixture shaken 18 hours. CPG was filtered successively with dichloromethane, and washed triethylamine, and dichloromethane. The resulting CPG was then dried under vacuum, suspended in 15 ml piperidine and shaken 30 minutes. CPG was filtered off, washed thoroughly 25 with dichloromethane and again dried under vacuum. The extent loading (determined by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm) was approximately 27 μ mol/g. The product solid support was subsequently used to synthesize the oligomers.

Preparation of 5'-0-dimethoxytrityl-3'-0-[hexyl-N-(1-pyrene propyl carbonyl] amino] uridine-2'-0-(succinyl amino propyl) controlled pore glass.

The procedure of Example 12 is repeated, except that 5'-O-dimethoxytrityl-3'-O-[hexyl-N-(1-pyrene propyl carbonyl] amino] uridine is used.

EXAMPLE 14

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(5-thio 10 carbonyl-3,6-dipivolyl-fluorescein)amino]uridine.

Fluorescein isothiocyanate (Isomer I, available from Cal Biochem, La Jolla, CA) was treated with 12 equivalents of in Et₃N/THF to give di-O-pivaloyl pivaloyl chloride fluorescein isothiocyanate. This compound was purified in 15 silica gel column using 3:1 hexane:ethyl acetate. Nucleoside. 5'-O-(dimethoxytrityl)-2'-O-(hexylamino)uridine was condensed with dipivolyl fluorescein isothiocyanate in 5'-0-The resultant compound CH₂Cl₂/pyrimidine. (dimethoxytrityl) -2'-O-[hexyl-N-(5-thiocarbonyl-3,6-dipivolyl-20 fluorescein) amino] uridine is then purified by using 100% ethyl acetate, in a silica column.

EXAMPLE 15

Preparation of 5'-0-dimethoxytrityl-2'-0-[hexyl-N-(5-thiocarbonyl-3,6-di-pivalcyl fluorescein) amino] uridine-3'-0-25 (2-cyanoethyl, N-N-diisopropyl phosphoramidite.

5'-O-(dimethoxytrity1)-2'-O-[hexyl-N-(5-thiocarbonyl-3,6-dipivolyl fluorescein)amino]uridine (0.75 grams, 0.672 mmol) was dissolved in dry dichloromethane (20 mL). 2-Cyanoethyl N,N,N',N'-tetraisopropylphosphorodiamidite (700 μL, 30 2.2 mmol) and diisopropylamine tetrazolide were added to the mixture, which was stirred under argon for 16 hours. The reaction mixture was then concentrated in vacuo and the residue dissolved in dichloromethane (75 mL) followed by washing with an equal volume of saturated NaHCO₃. The aqueous 15 layer was washed with dichloromethane (50 mL) and the combined

organic layers washed with an equal volume of saturated NaCl. The aqueous layer was washed with dichloromethane (50 mL) and the combined organic layers dried over MgSO₄ and concentrated. The residue was chromatographed on a silica gel column, 5 eluting with a gradient of 25% ethyl acetate in hexanes to 100% ethyl acetate. The desired product (670 mg, 77% yield, R, 0.79 by TLC) eluted with 100% ethyl acetate.

EXAMPLE 16

preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(5-10 thiocarbonyl-3,6-di-pivaloyl fluorescein)amino]uridine-3'-0-(succinylaminopropyl) controlled pore glass.

Succinylated and capped aminopropyl controlled pore glass (CPG) is dried under vacuum for 3 hours immediately before use. CPG (0.3 grams) is added to 3 ml anhydrous 15 pyridine in a 50 ml round-bottom flask. DEC (0.12 grams, 0.63 mmol), TEA (25 μ l, distilled over CaH₂, DMAP (dimethyl amino pyridine) (0.005 grams, 0.04 mmol) and 5'-O-dimethoxytrityl-2'-O-[hexyl-N-(5-thiocarbonyl-3,6-di-pivaloyl fluorescein) amino] uridine (0.21 grams, 0.19 mmol) are added under argon 20 and the mixture shaken mechanically for 19 hours. nucleoside (0.025 grams) is added and the mixture shaken an additional 5.5 hours. Pentachlorophenol (0.045 grams, 0.17 mmol) is added and the mixture shaken 18 hours. filtered off and washed successively with dichloromethane, 25 triethylamine, and dichloromethane. CPG then is dried under vacuum, suspended in 15 mL piperidine and shaken 30 minutes. CPG is filtered off, washed thoroughly with dichloromethane, and again dried under vacuum. The extent of loading is then determined by spectrophotometric assay of dimethoxytrityl 30 cation in 0.3 M p-toluenesulfonic acid at 498 nm.

EXAMPLE 17

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(3-oxycarbonyl-cholesteryl)amino]uridine.

Nucleoside 5'-O-(dimethoxytrityl)-2'-O-[hexylamino]-35 uridine (3.85 g, 6.0 mmol) was dissolved in anhydrous

pyridine/dichloromethane 50/50 (v/v) (20 mL). Cholesteryl chloroformate (Fluka, 3.0 g, 6.68 mmol) was dissolved in anhydrous dichloromethane (20 ml) and added slowly under argon with a syringe to the stirring reaction mixture. The mixture 5 was stirred under argon at room temperature for 2 h after Residual DMF was which it was concentrated in vacuo. coevaporated with toluene. The residue was dissolved in dichloromethane (50 mL) and washed with an equal volume The aqueous layer was washed with saturated NaHCO. 10 dichloromethane and the combined organic extracts washed with an equal volume saturated NaCl. The aqueous layer was washed with dichloromethane and the combined organic layers dried over MgSO4 and concentrated. The residue was chromatographed on a silica gel column with a gradient of 25% ethyl acetate 15 in hexanes to 100% ethyl acetate. The desired product (3.78 g, 60%) eluted with 100% ethyl acetate (R, 0.41 by TLC).

EXAMPLE 18

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(3-oxy-carbonyl-cholesteryl)amino]uridine-3'-0-[2-cyanoethyl-N,N-di-20 isopropyl]phosphoramidite

5'-O-(dimethoxytrityl)-2'-O-[hexyl-N-(3-Nucleoside oxycarbonyl-cholesteryl)amino]uridine (3.44 g, 3.3 mmol) was dissolved in dry dichloromethane (75 mL). N, N, N'N'-tetraisopropylphosphorodiamidite (Sigma, 2.1 ml, 6.6 25 mmol) and diisopropylamine tetrazolide (0.29 g, 1.7 mmol) were added to the mixture, which was stirred under argon for 16 H. Dichloromethane (75 mL) was added to the solution, which was washed with an equal volume of saturated NaHCO3. The aqueous layer was washed with an equal volume of dichloromethane. The 30 aqueous layer was washed with dichloromethane (30 ml) and the combined organic layers washed with an equal volume of saturated NaCl. The aqueous layer was washed with dichloromethane (30 mL) and the combined organic layers dried over concentrated in vacuo. The residue and 35 chromatographed on a silica gel column with a gradient of 25% ethyl acetate in hexanes to 70% ethyl acetate. The desired product (3.35 g, 82% yield, $R_f=0.71$ by TLC in 50% ethyl acetate in hexanes) eluted with 50% ethyl acetate.

EXAMPLE 19

Preparation of 5'-0-(dimethoxytrity1)-2'-0-[hexy1-N-(3-5 oxycarbonyl-cholesteryl)amino]uridine-3'-0-(succinyl aminopropyl)-controlled pore glass.

Succinylated and capped controlled pore glass (0.3 grams) is added to 2.5 ml anhydrous pyridine in a 15 ml pear-DEC (0.07 grams, 0.36 mmol), TEA (100 μ l, shaped flask. 10 distilled over CaH₂), DMAP (0.002 grams, 0.016 mmol) and 5'-O-(dimethoxytrityl) -2'-0-[hexyl-N-(3-oxycarbonyl-cholesteryl) amino]uridine (0.25 grams, 0.23 mmol) are added under argon and the mixture shaken mechanically for 16 hours. More nucleoside (0.20 grams) is added and the mixture shaken an 15 additional 18 hours. Pentachlorophenol (0.03 grams, 0.11 mmol) is added and the mixture shaken 9 hours. CPG is filtered off and washed successively with dichloromethane, triethylamine, and dichloromethane. CPG is then dried under vacuum, suspended in 10 ml piperidine and shaken 15 minutes. 20 CPG is filtered off, washed thoroughly with dichloromethane The extent of loading is and again dried under vacuum. determined by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm as approximately 39 μ mol/g.

25 EXAMPLE 20

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(2,4-dinitrophenyl)amino]uridine.

5'-O-(dimethoxytrityl)-2'-O-(hexylamino)uridine (0.88 grams, 1.37 mmol) was dissolved in methanol (20 mL). 2,4-30 Dinitrofluorobenzene (DNFB, 0.25 grams, 1.37 mmol) was added and the mixture shaken on a mechanical shaker. The reaction was monitored by TLC. After 90 min, another 0.25 grams of DNFB was added and the reaction mixture shaken an additional 30 min, followed by addition of another 0.25 grams of DNFB.
35 After shaking 2.5 hours, the mixture was concentrated in vacuo

and chromatographed on a silica gel column, eluting with a gradient of 25% ethyl acetate in hexanes to 100% ethyl acetate. The desired product (0.51 grams, 46%) eluted with 100% ethyl acetate (R_f 0.85 by TLC).

5 EXAMPLE 21

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(2,4-dinitrophenyl) amino]uridine-3'-0-(2-cyanoethyl-N,N-diisopropyl) phosphoramidite.

5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(2,4-10 dinitrophenyl)amino]uridine (0.45 grams, 0.55 mmol) was dissolved in dry dichloromethane (12 mL). 2-Cyanoethyl N, N, N', N'-tetraisopropylphosphorodiamidite (380 μ L, 1.2 mmol) and diisopropylamine tetrazolide (0.041 grams, 0.024 mmol) were added to the mixture, which was stirred under argon for 15 16 hours. The reaction mixture was then concentrated in vacuo and the residue dissolved in dichloromethane (75 mL) followed by washing with an equal volume of saturated NaHCO3. aqueous layer was washed with dichloromethane (25 mL) and the combined organic layers washed with an equal volume of The aqueous layer was washed with 20 saturated NaCl. dichloromethane (25 mL) and the combined organic layers dried over MgSO4 and concentrated. The residue was chromatographed on a silica gel column, eluting with a gradient of 20% ethyl acetate in hexanes to 100% ethyl acetate. The desired product 25 (510 mg foam, 93% yield, $R_f \mid 0.70$ by TLC) eluted with 100% ethyl acetate. ³¹PNMR (CDCl₃): 150.56 and 150.82 ppm.

EXAMPLE 22

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(2,4-dinitrophenyl)amino]uridine-3'-0-(succinyl aminopropyl) 30 controlled pore glass.

Succinylated and capped controlled pore glass (0.3 grams) is added to 3 ml anhydrous pyridine in a 50 ml round-bottom flask. DEC (0.12 grams, mmol), TEA (25 μ l, distilled over CaH₂), DMAP (0.005 grams, 0.041 mmol) and 5'-O-(di-35 methoxytrityl)-2'-O-[hexyl-N-(2,4-dinitrophenyl)amino]uridine

(0.21 grams, 0.26 mmol) are added under argon and the mixture shaken mechanically for 19 hours. More nucleoside (0.025 grams) is added and the mixture shaken an additional 5.5 hours. Pentachlorophenol (0.045 grams, 0.16 mmol) is added 5 and the mixture shaken 18 hours. CPG is filtered off and washed successively with dichloromethane, triethylamine, and dichloromethane. CPG then is dried under vacuum, suspended in 15 ml piperidine and shaken for 15 minutes. CPG is filtered off, washed thoroughly with dichloromethane, and 10 again dried under vacuum. The extent of loading is determined by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm, as approximately 29 μmol/gm.

EXAMPLE 23

15 Preparation of 5'-O- (dimethoxytrityl)-2'-O- [hexyl-N- (Nα-Nimid-Di-FMOC-L-Histidyl) amino] uridine.

Nucleoside 5'-O-(dimethoxytrityl)-2'-O-(hexylamino)uridine (0.97 g, 1.51 mmol) was dissolved in dichloromethane
(25 mL) and cooled to 0°C in an ice bath. Nα,Nimid-Di-FMOC-L20 histine pentafluorophenyl ester (2.4 g, 3.1 mmol, purchased
from Sigma) and 1-hydroxybenzotriazole (0.32 g, 0.24 mmol,
purchased from Fluka) were added to the stirred reaction
mixture stirred under argon. After 15 minutes, the ice bath
was removed and the mixture stirred under argon at room
25 temperature for 72 h. The mixture was concentrated in vacuo
and chromatographed on a silica gel column, eluting with a
gradient of 50% ethyl acetate in hexanes to 70% ethyl acetate
in hexanes. The desired product (0.53 g, 28%) eluted with 70%
ethyl acetate (R_f 0.53 by TLC in 100% ethyl acetate).

30 EXAMPLE 24

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(Nα-Nimid-Di-FMOC-L-histidyl)-amino]-uridine-3'-0-[2-cyanoethyl-N,N-disopropyl]phosphoramidite.

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(N α -Nimid-Di-FMOC-L-35 histidyl)amino]uridine (1.9 g, 1.6 mmol) is dissolved in dry

dichloromethane (20mL). 2-Cyanoethyl N,N,N',N'-tetraisopropylphosphorodiamidite (800 μL, 2.4 mmol) and diisopropylamine tetrazolide (0.090 grams, 0.52 mmol) are added to the
mixture, which is stirred under argon for 20 hours. The
5 reaction mixture then is concentrated in vacuo and the residue
dissolved in dichloromethane (75 mL). The solution is washed
with an equal volume of saturated NaHCO₃. The aqueous layer
is washed with dichloromethane (20 mL) and the combined
organic layers washed with an equal volume of saturated NaCl.
10 The aqueous layer is washed with dichloromethane (20 mL) and
the combined organic layers dried over MgSO₄ and concentrated.
The residue is chromatographed on a silica gel column, eluting
with a gradient of 50% ethyl acetate in hexanes to 100% ethyl
acetate. The desired product elutes with 100% ethyl acetate.

15 EXAMPLE 25

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(Nα-Nimid-Di-FMOC)-L-histidyl) amino] uridine-3'-0-[succinylaminopropyl] controlled pore glass.

Succinylated and capped controlled pore glass (dried 20 under vacuum for 3 hours immediately before use; 0.3 grams) is added to 3 ml anhydrous pyridine in a 50 ml round-bottom DEC (0.12 grams, 0.63 mmol), TEA (25 μ l, distilled over CaH₂), DMAP (0.005 grams, 0.04 mmol) and 5'-O-(dimethoxytrityl) -2'-0-[hexyl-N-(Na-Nimid-Di-FMOC)-L-histidyl)amino]-25 uridine (0.21 grams, 0.17 mmol) are added under argon and the mixture shaken mechanically for 19 hours. More nucleoside (0.025 grams) is added and the mixture shaken an additional Pentachlorophenol (0.045 grams, 0.17 mmol) is added and the mixture shaken 18 hours. CPG is filtered off 30 and washed successively with dichloromethane, triethylamine, and dichloromethane. CPG then is dried under vacuum, suspended in 15 ml piperidine and shaken 15 minutes. filtered off, washed thoroughly with dichloromethane and again dried under vacuum. The extent of loading is determined by 35 spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm. to be approximately 27 $\mu mol/g$.

EXAMPLE 26

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(Ω -methyl-5 polyethylene glycol-propionoyl)amino]uridine.

Nucleoside 5'-0-(dimethoxytrityl)-2'-0-[hexylamino]uridine, (1 g, 1.55 mmol) is dissolved in anhydrous DMF (15 1-Hydroxybenzotriazole (0.24 g, 1.75 mmol) polyethylene glycol-propionic acid-NHS-ester (1.23 g, 1.75 10 mmol) are added to the reaction mixture. The mixture is stirred under argon at room temperature for 2 hours after which it is concentrated in vacuo. Residual DMF coevaporated with toluene. The residue is dissolved in dichloromethane (50 mL) and then washed with an equal volume 15 saturated NaHCO3. The aqueous layer is washed with dichloromethane and the combined organic extracts washed with an equal volume saturated NaCl. The aqueous layer is washed with dichloromethane and the combined organic layers dried over MgSO4 and concentrated. The residue is chromatographed 20 on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product (1.08 g, 58%) eluted with 100% ethyl acetate.

EXAMPLE 27

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(Ω-methyl-25 polyethylene glycol-propionoyl)amino]uridine-3'-0-(2-cyano-ethoxy-N,N-diisopropyl)phosphoramidite.

5'-O-(Dimethoxytrityl)-2'-O-[hexyl-N-(Ω-methyl-polyethylene glycol-propionoyl)amino]uridine (1.04 grams, 0.87 mmol) is dissolved in dry dichloromethane (20mL). 2-Cyano-30 ethyl N,N,N',N'-tetraisopropylphosphorodiamidite (800 μL, 2.4 mmol) and diisopropylamine tetrazolide (0.090 grams, 0.52 mmol) are added to the mixture, which is stirred under argon for 20 hours. The reaction mixture then is concentrated in vacuo and the residue dissolved in dichloromethane (75 mL). 35 The solution is washed with an equal volume of saturated

NaHCO₃. The aqueous layer is washed with dichloromethane (20 mL) and the combined organic layers washed with an equal volume of saturated NaCl. The aqueous layer is washed with dichloromethane (20 mL) and the combined organic layers dried over MgSO₄ and concentrated. The residue is chromatographed on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product elutes with 100% ethyl acetate.

EXAMPLE 28

10 Preparation of 5'-0-(dimethoxytrityl)-2'-0-[hexyl-N-(Ω-methyl-polyethylene glycol-propionoyl)amino]uridine-3'-0-(succinyl-aminopropyl) controlled pore glass.

Succinylated and capped controlled pore glass (CPG) is dried under vacuum for 3 hours immediately before use. 15 Controlled pore glass (0.3 grams) is added to 3 ml anhydrous pyridine in a 50 ml round-bottom flask. DEC (0.12 grams, 0.67 mmol), TEA (25 μ l, distilled over CaH₂), DMAP (0.005 grams, 5'-O-(dimethoxytrityl)-2'-O-[hexyl-N-(w-methyland polyethylene glycol-propionoyl)amino]uridine (0.21 grams, 20 0.175 mmol) are added under argon and the mixture shaken mechanically for 19 hours. More nucleoside (0.025 grams) is added and the mixture shaken an additional 5.5 hours. Pentachlorophenol (0.045 grams, 0.17 mmol) is added and the CPG is filtered off and washed mixture shaken 18 hours. triethylamine, dichloromethane, 25 successively with dichloromethane. CPG then is dried under vacuum, suspended in 15 ml piperidine, and shaken 15 minutes. CPG is filtered off, washed thoroughly with dichloromethane, and again dried The extent of loading is determined by under vacuum. 30 spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm. to be approximately 18 μ mol/g.

Preparation of Macrocycle derivatized nucleoside

5'-O-(dimethoxytrityl)-2'-O-(hexylamine) uridine is treated as per the procedure of Example 3 with the macrocycle 54-{1,4,8,11-tetraza-[tri-(trifluoroacetyl) cyclotetradec-1-yl]}methyl benzoic acid-N-hydroxy succinimide ester (prepared according to Simon Jones, et. al., Bioconjugate Chem. 1991, 2, 416) to yield the product.

EXAMPLE 30

10 Preparation of macrocycle derivatized uridine phosphoramidite

The nucleoside product of Example 29 is treated as per
the procedure of Example 4 to yield the product.

EXAMPLE 31

Preparation of CPG derivatized with macrocycle derivatized
15 nucleoside

The nucleoside product of Example 29 is treated as per the procedure of Example 5 to yield the product.

EXAMPLE 32

Preparation of 5'-0-(dimethoxyltrityl)-2'-0-(hexyl-N-(folate)-20 amino)uridine

5'-O-(Dimethoxytrityl)-2'-O-(hexylamine)uridine is treated as per the procedure of Example 3 with folic acid pentafluorophenyl ester (protected with an isobutyryl protecting group) to yield the product.

25 EXAMPLE 33

Preparation of 5'-0-(dimethoxyltrityl)-2'-0-[hexyl-N-(folate)-amino]uridine-3'-0-(2-cyanoethoxy-N,N-diisopropyl)phosphoramidite

The nucleoside product of Example 29 is treated as per 30 the procedure of Example 4 to yield the product.

Preparation of CPG derivatized with 5'-0-(dimethoxyltrityl)-2'-0-(hexyl-N-(folate)amino)uridine nucleoside

The nucleoside product of Example 32 is treated as per 5 the procedure of Example 5 to yield the product.

EXAMPLE 35

5'-0-(dimethoxytrityl)-2'-0-{hexyl-N-[2of Preparation methoxy-6-chloro-9 (Ω -amino-caproyl) acridine] amino uridine.

6,9-Dichloro-2-methoxyacridine (Aldrich, 10 g, 36 mmol) 10 and phenol (2.5 g) were placed together on a round-bottom flask with a stirring bar and to this 6-amino-hexanoic acid (9.3 q, 71 mmol) was added and the flask was heated to 100° (oil bath) for 2 hours. TLC (10% methanol in methylene chloride) showed complete disappearance of starting material.

15 The reaction mixture was cooled and poured into 200 mL of methanol. The product isolates out as a yellow solid (about This compound was then converted into 10 q). pentafluorophenyl ester.

5'-O-(Dimethoxytrityl)-2'-(hexylamino)uridine (0.5g, 20 0.78 mmol) is dissolved in anhydrous DMF (15 mL). 1-Hydroxybenzotriazole (0.16 grams, 1.17 mmol) and 2-methoxy-6-chloro-9- $(\Omega$ -caproyl-amino) acridine pentafluorophenyl ester (0.53 grams, 1.17 mmol) are added to the reaction mixture. mixture is stirred under argon at room temperature for 2 h, 25 after which it is concentrated in vacuo. Residual DMF is coevaporated with toluene. The residue is dissolved in dichloromethane (50 mL) and washed with an equal volume saturated NaHCO3. The aqueous layer is washed with dichloromethane and the combined organic extracts washed with 30 an equal volume saturated NaCl. The aqueous layer is washed with dichloromethane and the combined organic layers dried over MgSO4 and concentrated. The residue is chromatographed on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product

35 elutes with 100% ethyl acetate.

Preparation of 5'-0-(dimethoxytrityl)-2'-0-{hexyl-N-[2-methoxy-6-chloro-9-(Ω-amino-caproyl)acridine]amino}uridine-3'-0-(2-cyanoethyl-N-N-diisopropyl) phosphoramidite.

5'-O-Dimethoxytrityl-2'-O-{hexyl-N-[2-methoxy-6-chloro-9-(w-amino-caproyl) acridine] amino] uridine (0.80 grams, 0.77 mmol) is dissolved in dry dichloromethane (20mL). ethyl N, N, N', N'-tetraisopropylphosphorodiamidite (800 μ L, 2.4 mmol) and diisopropylamine tetrazolide (0.090 grams, 0.52 10 mmol) are added to the mixture, which is stirred under argon for 20 hours. The reaction mixture is then concentrated in vacuo and the residue dissolved in dichloromethane (75 mL). The solution is washed with an equal volume of saturated NaHCO3. The aqueous layer is washed with dichloromethane (20 15 mL) and the combined organic layers washed with an equal volume of saturated NaCl. The aqueous layer is washed with dichloromethane (20 mL) and the combined organic layers dried over MgSO₄ and concentrated. The residue is chromatographed on a silica gel column, eluting with a gradient of 50% ethyl 20 acetate in hexanes to 92% ethyl acetate. The desired product elutes with 100% ethyl acetate.

EXAMPLE 37

Preparation of 5'-O-(dimethoxytrity1)-2'-O-{hexy1-N-[2-methoxy-6-chloro-9-(Ω-aminocaproy1)acridine]amino}uridine-3'-25 O-(succinyl aminopropyl) controlled pore glass.

Succinylated and capped controlled pore glass (0.3 grams) is added to 3 ml anhydrous pyridine in a 50 ml round-bottom flask. DEC (0.12 grams, 0.67 mmol), TEA (25 μ l, distilled over CaH₂), DMAP (0.005 grams, 0.04 mmol) and 5'-O-30 dimethoxytrityl-2'-O-{hexyl-N-[2-methoxy-6-chloro-9-(Ω -aminocaproyl)acridine]amino}uridine (0.21 grams, 0.17 mmol) are added under argon and the mixture shaken mechanically for 19 hours. More nucleoside (0.025 grams) is added and the mixture shaken an additional 5.5 hours. Pentachlorophenol 35 (0.045 grams, 0.17 mmol) is added and the mixture shaken 18 hours. CPG is filtered off and washed successively with

dichloromethane, triethylamine, and dichloromethane. CPG is then dried under vacuum, suspended in 15 ml piperidine and shaken 15 minutes. CPG is filtered off, washed thoroughly with dichloromethane and again dried under vacuum. The extent of loading is determined by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm. to be approximately 27 \(\mu\text{mmol/g}.\)

EXAMPLE 38

Preparation of 5'-0-(dimethoxytrityl)-2'-0-[(hexyl-N,N-10 dimethyl)amino]uridine.

5'-0-(dimethoxytrityl)-2'-0-(hexylamino)uridine (0.19 grams, 0.29 mmol) is dissolved in 4 ml methanol. acetate pH 4.0 (2 ml), sodium cyanoborohydride (0.02 grams, 0.3 mmol) and 37% formaldehyde in water (300 μ l) are added to 15 the reaction mixture, which is stirred 2 hours, after which it is concentrated in vacuo. The residue is dissolved in dichloromethane (50 mL) and washed with an equal volume The aqueous layer is washed with saturated NaHCO3. dichloromethane and the combined organic extracts washed with 20 an equal volume saturated NaCl. The aqueous layer is washed with dichloromethane and the combined organic layers dried over MgSO, and concentrated. The residue is chromatographed on a silica gel column, eluting with a gradient of 50% ethyl acetate in hexanes to 100% ethyl acetate. The desired product 25 (0.15 grams, 80%) elutes with 10% Methanol-90% ethyl acetate.

EXAMPLE 39

Oligonucleotides Having 3'-Alkylamino Group

3'-O-Hexyl-(N-phthalimido)-aminouridine-CPG, i.e. the 5'-O-dimethoxytrityl-3'-O-[hexyl-(Ω-N-phthalimido amino)]-uri-30 dine-2'-O-(succinyl-aminopropyl) controlled pore glass from Example 5, was used to synthesize the following oligonucleotides:

Oligomer 49: GAC U*

Oligomer 50: GCC TTT CGC GAC CCA ACA CU

35 Oligomer 51: GCC TTT CGC GAC CCA ACA CU*

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wherein "*" denotes the 3'-0 hexylamino-modified nucleoside.

Standard commercial phosphoramidites were used with the synthesis cycle times specified by the manufacturer in a 380B ABI instrument (Applied Biosystems).

Oligomer 49 was used for structural proof of 3'-O-alkylamine-bearing oligonucleotides at the 3'-terminal end. It showed the expected three ³¹P NMR signals (-0.5 ppm, -0.25 ppm, -0,2 ppm) and seven lines in the trace aromatic base region in ¹H NMR its spectrum.

Oligomer 51 was used to demonstrate the nuclease resistance offered by this the alkylamino group and also for further conjugation. The oligomer was treated with pyrenebutyric acid-N-hydroxy succinimide ester in 0.2 M NaHCO₃ buffer/DMF. The product, Conjugate 1, was purified by HPLC 15 and size exclusion methods. HPLC retention times (eluting with a gradient of 5% CH₃CN for 10 minutes then 5%-40% CH₃CN for 50 minutes) were as follows:

		1	Retention Time (min.)
20	Oligomer 50	1	25.99
	Oligomer 51	l	25.91
	Conjugate 1	1	49.35

The nuclease stability of Oligomer 51 and the conjugate were tested against Oligomer 50 in HeLa cytoplasmic/nuclear 25 extracts. The cell extract was diluted 1.4 times. The final concentration of oligonucleotide was 20 μ M. The half lives of the oligonucleotides were as follows:

		t _{1/2} (hours)
Oligomer 50		1.0
	1	3.5 3.6
	Oligomer 51	

The half life of phosphodiester Oligomer 50 increased 3-4 times by simple modification at the 3'-end with the 35 hexylamino group, by itself, or by further conjugation.

15

EXAMPLE 40

Oligonucleotides Having 2'-O-Alkylamino Group

A. The phosphoramidite from Example 4, 5'-O-(dimethoxytrityl)-2'-O-[hexyl-(Ω-N-phthalimido)amino]-uridine-53'-O-[(2-cyanoethyl)-N,N-diisopropyl]phosphoramidite was made as a 0.2 M solution in anhydrous CH₃CN and used to synthesize the following oligonucleotides in an ABI DNA synthesizer, model 380 B. During the modified amidite coupling, the reaction time was increased to 10 minutes. A coupling 10 efficiency of approximately 90% was observed. After deprotection with concentrated ammonium hydroxide (55°C, 16 hours) the oligonucleotides were purified by reverse phase HPLC and desalting column (Sephadex G-25).

Oligomer 52 (SEQ ID NO:14): GCGTGU*CTGCG Oligomer 53: GAU*CT

B. GCGTGTU'CTGCG where U' is 2'-0-[hexyl-N-(1-pyrene-propyl-carbonyl)amino uridine, Conjugate 2 (Oligomer 52 -pyrene butyrate conjugate).

To 20 O.D. of Oligomer 52 in 200 μ L of 0.2 M NaHCO₃ 20 buffer, 5 ml of pyrene-butyric acid-N-hydroxy succinimide ester in an Eppendorf tube was added followed by 200 μ L of DMF. The tube was shaken overnight. The reaction was purified by size exclusion and HPLC to yield 18 O.D. of product.

25 C. GCGTGTU"CTGCG where U" is 2'-O-[6-bromoacetamido-hex-1yl]-uridine, Conjugate 3 (Oligomer 52 - bromoacetate conjugate).

To 12 O.D. of Oligomer 52 in 100 μ L of 0.2 M NaHCO₃ buffer, 2 mg bromoacetic acid-NHS ester (N-hydroxy 30 succinimidyl bromoacetate) was added. After leaving the reaction to stand overnight, it was purified by size exclusion and HPLC to yield 7.5 O.D. of product.

- D. GCGTGTU^CTGCG where U^ is 2'-O-[hexyl-N-(polyethylene glycol)-propionoyl]amino uridine,
- 35 Conjugate 4 (Oligomer 52 PEG conjugate).

To 24 O.D. of Oligomer 52 in 200 μL of 0.2 M NaHCO₃ buffer, 20 mg of Polyethylene glycol propionic acid-N-hydroxy

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succinimide ester was added. The reaction was mechanically shaken overnight and purified by Sephadex G-25 size exclusion and chromatography to yield 22 O.D. of product.

HPLC retention times (eluting with a gradient of 5% 5 CH₃CN for 10 minutes then 5%-40% CH₃CN for 50 minutes in a C-18 Delta-Pak reverse phase column) were as follows:

		4 1	Retention Time (min.)
		1	
	Oligomer 52	1	24.05
10	Conjugate 2	l I	40.80
	Conjugate 3	i	26.04
	Conjugate 4	E .	55.58

Changes in T_m due to pyrene conjugation were evaluated against both DNA and RNA. T_m was measured in 100 mM Na $^{+}$, 10 15 nM phosphate, 0.1 mM EDTA, pH 7 at 4 μ M strand concentration. The results were as follows:

		T_m v. DNA (°C)	T_m v. RNA (°C)
	Oligomer 52	50.9	55.5
	Conjugate 2	55.3	55.5
20		(4.4)	(0.0)

The values in parentheses are changes in T_m compared to amino linker in oligomer 52 as a control.

EXAMPLE 41

Oligonucleotide synthesis using 2'-O-hexylamino(pyrene-25 butyrate)uridine phosphoramidite.

The amidite 5'-O-(dimethoxytrityl)-2'-O-[hexyl-N-(1-pyrene propyl carbonyl)amino]uridine-3'-O-(2-cyanoethyl-N,N-diisopropyl) phosphoramidite (0.2M in anhydrous acetonitrile) was used to synthesize the following oligomers, both for NMR 30 studies:

Oligomer 54: GAU CT

Oligomer 55: GCC GU*G TCG

(U'=2'-O-modified phosphoramidite)

These oligomers were purified trityl-on reverse-phase 35 HPLC, detritylated in 80% acetic acid for one hour and then repurified by RP-HPLC and desalted by size-exclusion

chromatography. NMR analysis showed the presence of pyrene peaks.

EXAMPLE 42

Oligonucleotide synthesis using 2'-O-hexylamino(dinitro-5 phenyl)uridine phosphoramidite.

The amidite 5'-O-(dimethoxytrityl)-2'-O-[hexyl-N-(2,4-dinitrophenyl) amino] uridine-3'-O-(2-cyanoethyl-N,N,-diiso-propyl) phosphoramidite (0.18 M in anhydrous acetonitrile) was used to synthesize oligonucleotides, Oligomers 56 to 63. All 10 are analogues of an ICAM antisense sequence. These oligomers were purified trityl-on by RP-HPLC (Waters Delta-Pak C18 column, 300 Å, 7.8 mm x 30 cm, linear 50-min gradient of 5-60% acetonitrile in 0.05 M TEAA pH 7.3), detritylated in 80% acetic acid for one hour and then purified by RP-HPLC and 15 desalted by size-exclusion chromatography. Data are summarized below:

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	Backbone	Total (O.D.)	Retention Time (min)
Oligomer 56: 5 GAU*CT	P=O	40	39.16
Oligomer 57: (SEQ ID NO:15) U'GG GAG CCA TAG CGA GGC#	P=S	64	39.19
Oligomer 58: 10 U'GG GAG CCA TAG CGA GGC	P=S	45	39.21
Oligomer 59: U'GG GAG CCA TAG CGA GGC	P=O	60	37.68
Oligomer 60: U'GG GAG CCA U'AG CGA GGC	P=O	69	38.58
15 Oligomer 61: TGG GAG CCA U*AG CGA GGC (SEQ ID NO:16)	P=O	86	32.38
Oligomer 62: U'CT GAG TAG CAG AGG AGC TC 20 (SEQ ID NO:17)	P=0	34	35.76
Oligomer 63: U'GG GAG CCA U'AG CGA GGC#	P=S	72	43.37

#=Non-nucleoside 6-carbon amino linker (Glen Research) and Bold indicates nucleotides having 2'-O-methyl substitutions

25 EXAMPLE 43

Oligonucleotide synthesis using 2'-O-[hexylamino-(cholesterol)]uridine phosphoramidite

The amidite 5'-O-dimethoxytrityl-2'-O-[hexyl-N-(3-oxycarbonyl-cholesteryl)amino]uridine-3'-O-[2-cyanoethyl-N,N,-30 diisopropyl]-phosphoramidite (0.2M in anhydrous acetonitrile/dichloromethane 2:1 v/v) was used to synthesize Oligomers 67-74. These oligomers are purified trityl-on by reverse-phase HPLC (Waters Delta-Pak C₁₈, 300Å, 7.8 mm x 30 cm, linear 55-min gradient of 5-80% acetonitrile in 0.05 M 35 TEAA pH 7.3), detritylated in 80% acetic acid for one hour and then repurified by RP-HPLC and desalted by size-exclusion chromatography. Data are summarized below:

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		Backbone	Target	Retention Time (min)
5	Oligomer 67: GAU'CT	P=O	NMR	52.73
	Oligomer 68: U'GG GAG CCA TAG CGA GGC	P=O	ICAM	49.64
10	Oligomer 69: U'GC CCA AGC TGG CAT CCG TC (SEQ ID NO:18)	P=S	ICAM	51.98
	Oligomer 70: U'GC GTT TGC TCT TCT TG (SEQ ID NO:19)	P=S C G	CMV	52.57
15	Oligomer 71: U'GC ATC CCC CAG GCC ACC AT (SEQ ID NO:20)	P=S	mseICAM	53.24
	Oligomer 72: U'CC CGC CTG TGA CAT GCA TT (SEQ ID NO:21)	P=S	Raf	53.95
20	Oligomer 73: GU'T CTC GCT GGT GAG TTT CA (SEQ ID NO:22)	P=S	PKCa	51.04
25	Oligomer 74: F1-UU*GG GAG CCA TAG CGA GG (SEQ ID NO:23)	1	ICAM	52.75
	Fl-U = U 2'-modified with i	Tuorescein	(see rxswbr	C 44).

EXAMPLE 44

Synthesis of oligonucleotides using 2'-0-[hexylamino-(fluorescein)] amidite.

The amidite 5'-O-dimethoxytrityl-2'-O-[hexyl-N-(5-thiocarbonyl-3,6-dipivolyl fluorescein)amino]uridine-3'-O-(cyanoethyl-N,N-diisopropyl phosphoramidite) (0.2 M in anhydrous acetonitrile) was used to synthesize Oligomer 74 (above) and Oligomers 75-82 on a 1 x 10⁵ (Oligomer 75) or 1 x 35 10² (remaining Oligomers) μmol scale. These oligomers are purified trityl-on by reverse phase HPLC (Waters Delta-Pak C₁₈, 300Å, 7.8 mm x 30 cm, linear gradient of acetonitrile in 0.05 M TEAA pH 7.3), detritylated in 80% acetic acid for one

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hour and then repurified by RP-HPLC and desalted by size-exclusion chromatography.

					1		Backbone	Target
5	Oligomer GAU'CT	75:			i 1		P=O	NM R
	Oligomer U'GG GAG		G CGA	GGC	1		P=O	ICAM
	Oligomer U'GC CCA		G CAT	CCG	TCA		P=S	ICAM
10	Oligomer U'GC CCA		G CAT	CCG	TCA	#	P=S	ICAM .
	Oligomer U'GC GTT		T TCT	тст	TGC	! G	P=S	CMV
15	Oligomer U'GC ATC		.G GCC	ACC	AT		P=S	mseICAM
	Oligomer U'GC ATC (U-CPG) (U-CPG	CCC CA	G GCC	ACC	A	nido U 6	P=S	mseICAM
				L		1		
20	Oligomer GU'T CTC		T GAG	TTT	CA	1	P=S	PKC
	Where U'	is U m	odifie	ed w	ith	fluoresce	in.	

EXAMPLE 45

1-Adamantane Acetic Acid Pentafluorophenyl Ester

1-Adamantane acetic acid (5 g, 25.77 mmol) was 25 dissolved in 50 mL of CH₂Cl₂ and to this suspension (5 pentafluorophenyl 27.16 mmol) g, Dicylohexylcarbodiimide (DCC) was then added (6 g, 29.12 mmol), at which point a highly exothermic reaction set in. The 30 reaction flask was saturated with argon, sealed and then shaken in a wrist action shaker. After 5 hours the solution was filtered in a sintered glass funnel to filter off the dicylohexylurea formed during the reaction. The methylene chloride solution was evaporated and the pale yellow solid 35 product (9g, 97% crude product yield) obtained was used in the next condensation step.

5'-O-Dimethoxytrity1-2'-O-[hexyl-N-(1-adamantane acetamido)-amino]uridine

5'-O-Dimethoxytrityl-2'-O-(hexylamino)uridine (2 g, 5 3.15 mmol) was dissolved in 20 mL of methylene chloride. Adamantane acetic acid pentafluorophenyl ester (1.8 g from the previous step, 5 mmol crude product) was added followed by 3 The reaction mixture was shaken for 4 mL of triethylamine. hours in a wrist action shaker. TLC analysis (5% CH3OH in 10 CH₂Cl₂) showed the product formation as revealed by a moving trityl positive spot. When the amine at the origin was completely consumed (5 hours) the reaction mixture was evaporated and applied to a silica column loaded with chloroform solvent (20cm x 5cm diameter). The column was 15 eluted with chloroform (1 liter) followed by 5% CHCl3 in CH,OH. The material which showed trityl positive spots in TLC was pooled together and evaporated to give 2.5 g of the nucleoside-adamantane conjugate (98% yield). 13C analysis showed the presence of both uridine and adamantane residues. 20 R_f= 0.21 in 5% CH₃OH in CH₂Cl₂.

EXAMPLE 47

4

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(1-adamantane acetamido)-amino]uridine-3'-O-(2-cyanoethyl-N,N'-diisopropyl)
Phosphoramidite

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(1-adamantane acet-25 amido) amino] uridine (2.4 g, 2.96 mmol) was dissolved in 20 mL of CH₂Cl₂. To this solution 200 mg of diisopropylaminotetrazolide (1.16 mmol) was added followed by 3.1 mL of 2cyanoethyl-N,N,N',N'-tetraisopropyl phosphoramidite (9.45 30 mmol) was added under argon atmosphere. The mixture was stirred under argon overnight after which time TLC (5% CH₃OH/CH₂Cl₂) indicated almost complete conversion nucleoside to its phosphoramidite. The reaction mixture was washed with saturated NaHCO3 followed by saturated NaCl 35 solution and then dried over anhydrous MgSO4. The solution was evaporated and the foamy residue was purified in a silica column using 5% CH3OH/CHCl3 elution system to give the

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purified phosphoramidite (2 g, 67% yield). ³¹P NMR: 150.741, 150.49 ppm.

EXAMPLE 48

20

Oligonucleotide Synthesis with Adamantane Modified Amidite

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(1-adamantane acetamido)amino]uridine phosphoramidite was dissolved in anhydrous CH₃CN (100 mg of amidite for each 1 mL of solvent to give 0.1 M solution of the amidite). For the novel amidite the coupling time was extended by increasing the amidite and 10 delivery time to 7.5 minutes and the waiting time to 6.0, thus a total of 13.5 minutes were used.

The oligonucleotides were deprotected and purified using the procedure described in Example 2. Table I summarizes the different oligonucleotides along with their 15 HPLC retention times.

EXAMPLE 49

Table of Oligonucleotides Synthesized with their HPLC Retention Times - Incorporation of 2'-O-(CH₂)₆-NHCO-O-adamantylmethyl Uridine using amidite:

SEQUENCE	BB	RT (min)	TARGET
U*T	P=O	a	NMR
GAU*CT	P=O	29.3	NMR ·
U*GG GAG CCA TAG CGA GGC	P=O	a	ICAM
TGG GAG CCA U*AG CGA GGC	P=O	26.8	ICAM
U*GC ATC CCC CAG GCC ACC	P=S	31.3/ 32.0	mICAM
U*GC ATC CCC CAG GCC ACC	P=S	31.9	mICAM
U* CAG TGC CTG CGC CGC GCT CG (SEQ ID NO:27)	P=S	31.5/ 32.0	Ki-ras
U* CC GTC ATC GCT CCT CAG GG (SEQ ID NO:28)	P=S	31.6/ 32.1	H-ras
U*CC CGC CTG TGA CAT GCA TT a not determined	P=S	31.8/ 32.3	Raf

HPLC conditions: Waters DeltaPak C-18 Delta Pak reverse-phase 15m 300 Å column, 3.9x 300 mm size equipped with C-18 guard column; Solvent A: 50 mM TEAA, pH 7.0.; Solvent B: acetonitrile; Gradient: 0-10 min., 5% acetonitrile; 10-60 5 min., 5% to 60% acetonitrile linear increase.

EXAMPLE 50

cis-11-Eicosenoic Acid Pentafluorophenyl Ester

cis-11-Eicosenoic acid (Aldrich, 2 g, 6.44 mmol) was dissolved in 50 mL of THF and to this suspension 1.2 g of 10 pentafluorophenyl (6.5 mmol) was added followed by 1.33 g of DCC. The suspension was saturated with argon, sealed with a rubber septum and then shaken in a wrist action shaker. After 4 hours a clear pale yellow solution results with creamy white precipitate. The precipitated dicyclohexylurea was filtered 15 off, washed with methylene chloride and the combined organic solution was evaporated to give the desired product as a waxy solid (3.25 g, crude product yield). This solid was used in the next condensation step without any further purification.

EXAMPLE 51

20 5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(cis-11-eicosenoicamido)-amino]uridine

5'-O-Dimethoxytrityl-2'+O-(hexylamino)uridine (2 g, 3.15 mmol) was dissolved in 20 mL of methylene chloride. cis-11-Eicosenoic acid pentafluorophenyl ester (2.5 g from the previous step, 5 mmol crude product) was added followed by 3 mL of pyridine. The reaction mixture was shaken for 4 hours in a wrist action shaker. TLC analysis (5% CH₃OH in CH₂Cl₂) showed the product formation as revealed by a trityl positive spot above the origin. When the amine at the origin was completely consumed (5 hours) the reaction mixture was evaporated and applied to a silica column loaded with chloroform solvent (20 cm x 5 cm diameter). The column was eluted with chloroform (1 liter) followed by 5% CHCl₃ in CH₃OH. The material which showed trityl positive spots in TLC was pooled together and evaporated to give 2.72 g of the

nucleoside-eicosenoic acid conjugate (93% yield). 13 C analysis showed the presence of both uridine and cis-11-eicosenoic acid residues. $R_f=0.29$ in 5% CH_3OH in CH_2Cl_2

EXAMPLE 52

5 5'-O-Dimethoxtrityl-2'-O-[hexyl-N-(cis-11-eicosenoic amido) amino]uridine-3'-O-(2-cyanoethyl-N,N'-diisopropyl) Phosphoramidite

The eicosenoic acid-uridine nucleoside from the previous step (2.5 g, 2.7 mmol) was dissolved in 20 mL of CH₂Cl₂. 10 this solution 200 mg of diisopropylaminotetrazolide (1.16 mmol) was added followed by 2.0 mL of 2-cyanoethyl-N,N,N',N'tetraisopropyl phosphoramidite (6.1 mmol) was added under The mixture was stirred under argon argon atmosphere. overnight after which time TLC (5% CH₃OH/CH₂Cl₂) indicated 15 almost complete conversion of nucleoside to its phosphor-The reaction mixture was washed with saturated NaHCO₃ followed by saturated NaCl solution and then dried over anhydrous MgSO₄. The solution was evaporated and the foamy residue was purified in a silica column using 5% CH3OH/CHCl3 20 elution system to give the purified phosphoramidite (2 g, 67% yield). As the phosphoramidite was contaminated with the Hphosphorate from the phosphitylating reagent, repurified in a silica column, by eluting with 8:2 EtOAc: Hexane to give 1.8 g (60% yield) of the phosphoramidite. 31P 25 NMR: 150.74 and 150.56 ppm.

EXAMPLE 53

Oligonucleotide Synthesis with cis-11-eicosenoic acid modified amidite

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(cis-11-eicosenoic amido) amino] uridine phosphoramidite was dissolved in anhydrous CH₃CN (100 mg of amidite for each 1 mL of solvent to give 0.09 M solution of the amidite). For the novel amidite the coupling time was extended by increasing the amidite and delivery time to 7.5 minutes and the waiting time to 6.0, thus 35 a total of 13.5 minutes were used.

The oligonucleotides were deprotected and purified using the procedure described in Example 2. Table II summarizes the different oligonucleotides along with their HPLC retention times.

5 EXAMPLE 54
Table of Incorporation of 2'-0-(CH₂)₆-NHCO-Cis-11-Eicosenoic
U using amidite:

SEQUENCE	BB	RT (min)	TARGET
U*T	P=O	a	NMR
GAU*CT	P=O	a	NMR
U*GG GAG CCA TAG CGA GGC	P=O	a	ICAM
TGG GAG CCA U*AG CGA GGC	P=O	17.8	ICAM
U*GC ATC CCC CAG GCC ACC	P=S	55.5/ 56.4	mICAM
U*GC ATC CCC CAG GCC ACC	P=S	56.3/ 57.3	mICAM
U* CAG TGC CTG CGC CGC GCT CG	P=S	a	Ki-ras
U* CC GTC ATC GCT CCT CAG GG	P=S	57.0/ 57.8	H-ras
U*CC CGC CTG TGA CAT GCA	P=S	57.1/ 57.9	Raf

a not determined

HPLC conditions: Waters DeltaPak C-18 Delta Pak reverse10 phase15m 300 Å column, 3.9x 300 mm size equipped with C-18
guard column; Solvent A: 50mM TEAA, pH 7.0.; Solvent B:
acetonitrile; Gradient: 0-10 min., 5% acetonitrile; 10-60
min., 5% to 60% acetonitrile linear increase.

EXAMPLE 55

15 5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(N-biotinyl-6-aminocap-royl)amino]uridine

5'-O-Dimethoxytrityl-2'-O-(hexylamino)uridine (1.05 g, 1.66 mmol) was dissolved in 20 mL of methylene chloride. N-Biotinyl-6-amino caproic acid (0.75 g, Fluka, 1.65 mmol) was

added followed by 1 mL of pyridine. The reaction mixture was shaken for 4 hours in a wrist action shaker. The white suspension of the biotin reagent completely dissolved into solution by this time, TLC analysis (5% CH₃OH in CH₂Cl₂) showed 5 the product formation as revealed by an upper trityl positive spot above the origin. When the amine at the origin was completely consumed (5 hours) the reaction mixture was evaporated and applied to a silica column loaded with chloroform solvent (20 cm x 5 cm diameter). The column was 10 eluted with chloroform (1 liter) followed by 5% CHCl, in CH3OH (1 liter), then followed by 10% CH₃OH in CHCl₃. The material, which elutes out with 10% CH,OH, showed trityl positive spots in TLC was pooled together and evaporated to give 1.53 g of the nucleoside-biotin conjugate (98% yield). 13C analysis 15 showed the presence of both uridine and biotin residues. $R_{\rm f}$ = 0.214 in 10% CH₃OH in CH₂Cl_{2.}

EXAMPLE 56

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(N-biotinyl-6-aminocap-royl)amino]uridine 3'-O- (2-cyanoethyl-N,N'-diisopropyl) Phos-20 phoramidite

The biotin-uridine nucleoside from the previous step was dissolved in 20 mL of CH₂Cl₂. To this solution 100 mg of diisopropylaminotetrazolide (.58 mmol) was added followed by 1.0 mL of 2-cyanoethyl-N,N,N',N'-tetraisopropyl phosphoramidite (3.05 mmol) was added under argon atmosphere. The mixture was stirred under argon overnight after which time TLC (5% CH₃OH/CH₂Cl₂) indicates almost complete conversion of nucleoside to its phosphoramidite. The reaction mixture was washed with saturated NaHCO₃ followed by saturated NaCl solution and then dried over anhydrous MgSO₄. The solution was evaporated and the foamy residue was purified in a silica column using 100% ethyl acetate elution system to give the purified phosphoramidite.

5-O-Dimethoxytrityl-3'-O-[hexyl-N-(N-biotinyl-6-aminocaproyl)-aminoluridine

5'-O-Dimethoxytrityl-3'-O-(hexylamino)uridine (0.53 g, 5 0.83 mmol) was dissolved in 10 mL of methylene chloride. Biotinyl-6-aminocaproic acid (0.75 g, Fluka, 1.65 mmol) was added followed by 0.5 mL of pyridine. The reaction mixture was shaken for 4 hours in a wrist action shaker. suspension of the biotin reagent was completely dissolved into 10 solution by this time. TLC analysis (10% CH₃OH in CH₂Cl₂) showed the product formation as revealed by an upper trityl positive spot above the origin. When the amine at the origin was completely consumed (5 hours) the reaction mixture was evaporated and applied to a silica column loaded with 15 chloroform solvent (20 cm \times 5 cm diameter). The column was eluted with chloroform (1 liter) followed by 5% CHCl, in CH3OH The material, which elutes out with 10% CH3OH, (1 liter). showed trityl positive spots in TLC was pooled together and evaporated to give 0.6 g of the nucleoside-biotin conjugate 20 and biotin residues. R_f= 0.21 in 10% CH₃OH in CH₂Cl₂.

EXAMPLE 58

5'-O-Dimethoxytrityl-3'-O-[hexyl-N-(N-biotinyl-6-aminocaproyl) amino] uridine-2'-O-(succinylaminopropyl) Controlled Pore Glass

Succinylated/capped aminopropyl controlled pore glass was dries under vacuum for 3 hours immediately before use. A portion (0.3 g, 24 µmol) was added to 3 mL anhydrous pyridine in a 50 mL round-bottom flask. DEC (0.12 g, 0.63 mmol), TEA (25 µL, distilled over CaH₂). DMAP (0.005 g, 40 µmol) and 5′-30 O-(dimethoxytrityl)-3′-O-[hexyl-N-(N-biotinyl-6-aminocaproyl) amino]uridine (0.22 g, 0.22 mmol) were added under argon and the mixture shaken mechanically for 19 hours. More nucleoside (0.025 g) was added and the mixture shaken for an additional 5.5 hours. Pentachlorophenol (0.045 g, 0.168 mmol) was add 35 and the mixture shaken for 18 hours. CPG was filtered off and washed successively with dichloromethane, triethylamine, and

dichloromethane. The resulting CPG was then dried under vacuum, suspended in 15 mL piperidine and shaken 30 minutes. CPG was filtered off, washed thoroughly with dichloromethane and again dried under vacuum. The extent of loading (determined by spectrophotometric assay of dimethoxytrityl cation in 0.3 M p-toluenesulfonic acid at 498 nm) was approximately 36 μ mol/g. The product solid support was subsequently used to synthesize the oligomers, containing biotin at the 3' nucleotide unit.

10 EXAMPLE 59

Oligonucleotide Synthesis with Biotin Modified Amidite and Control Pore Glass

5'-O-Dimethoxytrityl-3'-O-[hexyl-N-(N-biotinyl-6-amino-caproyl) amino] uridine nucleoside phosphoramidite was dissolved 15 in anhydrous CH₃CN (100 mg of amidite for each 1 mL of solvent to give 0.08-0.09 M solution of the amidite). For the novel amidite the coupling time was extended by increasing the amidite and delivery time to 7.5 minutes and the waiting time to 6.0, thus a total of 13.5 minutes were used.

The oligonucleotides were deprotected and purified using the procedure described in Example 2. The controlled pore glass containing biotin was used to synthesize 3'-biotin containing oligonucleotides.

EXAMPLE 60

25 1-2-Di-O-hexadecyl-rac-glycerol-succinimidyl-carbonate

1,2-Di-O-hexadecyl-rac-glycerol (2g, 3.69 mmol) dissolved in 3 mL of CH₂Cl₂ and to this solution disuccinimidyl carbonate (DSC) (1.42 g, 5.54 mmol) was added. Triethylamine was then added (2 mL, 14.3 mmol), followed by 30 20 mL of anhydrous acetonitrile. The reaction flask was saturated with argon, sealed and then shaken in a wrist action After 8 hours the reaction mixture was evaporated. The residue was dissolved in CH2Cl2 (100 mL), washed with saturated NaHCO, solution followed by saturated NaCl solution 35 and the organic layer was dried over anhydrous MgSO₄.

solid residue (3.5 g) was used in the next step to condense with the amine. ¹³C NMR confirms the complete conversion of 1,2-di-O-hexadecyl-rac- glycerol to the corresponding succinimidyl carbonate.

5 EXAMPLE 61

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(carbonoyl 1,2-di-O-hexa-decyl-rac-glycerol)amino]uridine

5'-O-Dimethoxytrityl-2'-O-(hexylamino)uridine (2 g, 3.15 mmol) was dissolved in 20 mL of methylene chloride. 1,2-Di-O-10 hexadecyl-rac-glycerol succinimidyl carbonate (3.0 g from the previous step, 5 mmol crude product) was added followed by 3 mL of pyridine. The reaction mixture was shaken for 4 hours in a wrist action shaker. TLC analysis (5% CH₃OH in CH₂Cl₂) showed the product formation as revealed by a moving trityl 15 positive spot. When the amine at the origin was completely consumed (5 hours) the reaction mixture was evaporated and applied to a silica column loaded with chloroform solvent (20 cm x 5 cm diameter). The column was eluted with chloroform (1 liter) followed by 5% CHCl3 in CH3OH. The material which 20 showed trityl positive spots in TLC was pooled together and evaporated to give 3.0 g of the nucleoside-lipid conjugate (98% yield). 13C analysis showed the presence of both uridine and lipid residues.

EXAMPLE 62

25 5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(carbonoyl1,2-di-O-hexa-decyl-rac-glycerol-succinimidyl-carbonate]-3'-O-(2-cyanoethyl-N,N'-diisopropyl) Phosphoramidite

The lipid-uridine nucleoside from the previous step was dissolved in 20 mL of CH_2Cl_2 . To this solution 200 mg of 30 disopropylaminotetrazolide (1.16 mmol) was added followed by 1.55 mL of 2-cyanoethyl-N,N,N',N'-tetraisopropyl phosphoramidite (4.75 mmol) was added under argon atmosphere. The mixture was stirred under argon overnight after which time TLC (5% CH_3OH/CH_2Cl_2) indicated almost complete conversion of 35 nucleoside to its phosphoramidite. The reaction mixture was

washed with saturated NaHCO₃ followed by saturated NaCl solution and then dried over anhydrous MgSO₄. The solution was evaporated and the foamy residue was purified in a silica column using 5% CH₃OH/CHCl₃ elution system to give the purified phosphoramidite.

EXAMPLE 63

Oligonucleotide Synthesis with Lipid Modified Amidite

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(carbonoyl-1,2-di-O-hexadecyl-rac-glycerol)amino]uridine phosphoramidite was 10 dissolved in anhydrous CH₃CN (100 mg of amidite for each 1 mL of solvent to give 0.08-0.09 M solution of the amidite). For the novel amidite the coupling time was extended by increasing the amidite and delivery time to 7.5 minutes and the waiting time to 6.0, thus a total of 13.5 minutes were used. The 15 oligonucleotides were deprotected and purified using the procedure described in Example 2.

EXAMPLE 64

Oligonucleotide Synthesis

5'-O-Dimethoxytrityl-2'-O-[hexyl-N-(1-pyrene butyroyl)
20 amino]uridine phosphoramidite was dissolved in anhydrous CH₃CN
(100 mg of amidite for each 1 mL of solvent to give 0.08-0.09
M solution of the amidite). For the novel amidite the coupling time was extended by increasing the amidite and delivery time to 7.5 minutes and the waiting time to 6.0
25 minutes, hence using a total of 13.5 minutes.

The oligonucleotides were deprotected and purified using the procedure described in Example 2. The Table below summarizes the different oligonucleotides (ICAM, ras, raf and PKC-a oligonucleotides).

Table of Incorporation of 2'-O-(CH₂)₆-NHCO-butyryl pyrene U using amidite

SEQUENCE	BB	RT (min)	TARGET
U*T	P=O	40.66	NMR
U*GG GAG CCA TAG CGA GGC	P=O	a	ICAM
TGG GAG CCA U*AG CGA GGC	P=O	a	ICAM
U*GC ATC CCC CAG GCC ACC	P=S	30.00, 30.05	mICAM
U*GC ATC CCC CAG GCC ACC AT-3'-aminolink	P=S	30.29, 31.25	mICAM
U* CAG TGC CTG CGC CGC GCT	P=S	30.47, 31.32	Ki-ras
U* CC GTC ATC GCT CCT CAG	P=S	30.90, 31.66	H-ras
U*CC CGC CTG TGA CAT GCA	P=S	31.54/ 32.19	Raf
U*GT TCT CGC TGG TGA GTT TCA a not determined	P=S	30.96/ 31.86	PKC-a

HPLC conditions: Waters DeltaPak C-4 Delta Pak reverse-5 phase15m 300 Å column, 3.9x 300 mm size equipped with C-18 guard column; Solvent A: 50mM TEAA, pH 7.0.; Solvent B: acetonitrile; Gradient: 0-10 min., 5% acetonitrile; 10-60 min., 5% to 60% acetonitrile linear increase.

EXAMPLE 65

10 Incorporation of 2'-O-(CH₂)₆-NHCO-cholesterol Uridine at 5'-End of Oligonucleotides Using Amidites

Synthesis:

Oligonucleotides were synthesized on an Expedite Nucleic Acid Synthesis System (Millipore Corp., Bedford, MA). Two 1-15 μ mol syntheses or a single 15 μ mol synthesis was performed for each oligonucleotide. Standard DNA synthesis conditions were used except during coupling of the modified amidite which was accomplished as follows: The novel amidite was dissolved in CH₂Cl₂/CH₃CN 50/50 (0.08-0.1M). For 10 μ mol syntheses, a 6-min 20 wait step was added to the coupling cycle, which was repeated

once. For 1 μ mol phosphodiester syntheses, the coupling step was also extended for 6 min, with periodic addition of fresh amidite. For 1 μ mol phosphorothicate syntheses, the coupling step was extended for 13 min, with periodic addition of fresh 5 amidite.

Purification:

After standard cleavage from CPG and deprotection at | 55 °C for 16-24 hours), hydroxide (ammonium oligonucleotides were trityl-on purified by preparative HPLC 10 to remove failure sequences. HPLC conditions: Waters Delta-Pak C, column, 7.8 mm x 30 cm, 300 A, 15 u; 2.5 ml/min flow rate using a gradient of 5-80% AcCN in 50mM triethylammonium acetate pH 7.0 over 50 min. Following lyophilization, oligonucleotides were detritylated for 45 min in 80% acetic 15 acid and lyophilized again. Oligonucleotides were then further purified by size exclusion on G-25 columns to remove salts and the trityl group.

Analysis:

Except for 8002, oligonucleotides were analyzed by HPLC 20 using a gradient analogous to the one above and a Waters Delta-Pak C4 column (3.9 mm x 30 cm) with a flow rate of 1.5 ml/min over 50 min. 8002 was analyzed using a gradient of 5-60% AcCN. Purified phosphorothicate oligonucleotides appeared two peaks by HPLC, phosphodiesters as one peak. 25 Oligonucleotides were further analyzed by electrophoresis on 20% polyacrylamide denaturing (7M urea) gels. All were predominantly single bands of the appropriate size compared to standards. Gel analysis of 8012 (loaded without any shows a very high band, presumably the pretreatment) 30 tetrameric form of this oligonucleotide, plus a faint band where one would expect a monomeric 8mer+cholesterol to run. (The parent compound, 5320, is known to form a G-quartet structure.) Some oligonucleotides were also analyzed by capillary electrophoresis. Selected oligonucleotides have 35 been analyzed by electrospray mass spectrometry.

Results are shown in the following Table were * indicated site of selected modification.

	SEQUENCE	TYPE	HPLC RETENTION TIME (min)	TARGET
	U*GGGAGCCATAGCGAGGC	P=0	54.65	ICAM
	U*GCCCAAGCTGGCATCCGTCA	P=S	45.77, 46.44	ICAM
5	U*GCGTTTGCTCTTCTTCTTGCG	P=S	45.35, 46.03	CMV
-	U*GCATCCCCCAGGCCACCAT	P=S	56.74, 57.65	mICAM
	U*CCCGCCTGTGACATGCATT	P=S	46.69, 47.25	RAF
	U*TTAGGATTCGTGCTCATGG (SEQ ID NO:29)	P≃S	47.71, 48,32	HCV
10	U*GTTCTCGCTGGTGAGTTTCA (SEQ ID NO:30)	P=S	46.78, 47.58	PKC _a
	U*GGGAGCCATAGCGAGGC	P=S	not det'd	HIV
	U*TGGGGTT	P=S	48.46, 49.04	HIV

Example 66

15 Incorporation of 2'-O-(CH₂)₆-NHCO-cholesterol Uridine at nonterminal positions of oligonucleotides using amidite

Oligonucleotides incorporating $2'-0-(CH_2)_6-NHCO-$ cholesterol at nonterminal (not at either the 5' or the 3' terminus of the oligonucleotide) were synthesized at a 1 μ mol 20 syntheses scale, analyzed and purified as per the procedures of Example 65.

Results are shown in the following Table were * indicated site of selected modification.

25	SEQUENCE	TYPE	HPLC RETENTION TIME (min)	TARGET
	TGGGAGCCAU*AGCGAGGC	P=O	42.21	ICAM
	GU*TCTCGCTGGTGAGTTTCA	P=S	not det'd	PKC _«

Example 67

Incorporation of 3'-O-(CH₂)₆-NHCO-cholesterol Uridine at 3'end 30 of oligonucleotides using modified CPG

Cholesterol-modified CPG was packed into screw-top CPG columns (Glen Research). Oligonucleotides were synthesized on a 1- μ mol syntheses scale, analyzed and purified as in Example 65 except that standard DNA synthesis conditions were 35 used for all couplings.

Results are shown in the following Table were * indicated site of selected modification.

	SEQUENCE	TYPE	HPLC RETENTION TIME (min)	TARGET
	GCCTTTCGCGACCCAACACU* GCCTTTCGCGACCCAACACU*	P=S P=O	48.37 50.18	HCV HCV
5	TGCATCCCCCAGGCCACCAU* (SEQ ID NO:31)	P=O	47.01	mICAM
	GTGCTCATGGTGCACGGTCU* (SEQ ID NO:32)	P=S	49.97	HCV
	TGCATCCCCAGGCCACCAU*	P=S	≈ 59. 5	mICAM

10 Example 68

Incorporation of $2'-0-(CH_2)_6$ -NHCO-cholesterol Uridine at 3'end of oligonucleotides using modified CPG

Cholesterol-modified CPG was packed into screw-top CPG columns (Glen Research). Oligonucleotides were synthesized on a 1-µmol syntheses scale, 15 analyzed and purified as in Example 65 except that standard DNA synthesis conditions were used for all couplings.

Results are shown in the following Table were * indicated site of selected modification.

20	SEQUENCE	TYPE	HPLC RETENTION TIME (min)	TARGET
	TGCATCCCCCAGGCCACCAU*	P=0	47.03	ICAM

Example 69

Double Incorporation of Cholesterol by placement of (2'-0-(CH₂)₆-NHCO-cholesterol Uridine on 5' end and 3'-0-(CH₂)₆-NHCO-25 cholesterol Uridine on 3' end

Cholesterol-modified CPG was packed into screw-top CPG columns (Glen Research). Oligonucleotides were synthesized on a 1- μ mol syntheses scale, analyzed and purified as in Example 65 except that the extended coupling step for the 30 cholesterol amidite (see Example 65) was repeated once.

Results are shown in the following Table were * indicated site of selected modification.

`.	SEQUENCE	TYPE	HPLC RETENTION TIME (min)	TARGET
35	U*GCATCCCCCAGGCCACCAU*	P=S	54.31	ICAM
	(SEQ ID NO:33) U*GCATCCCCCAGGCCACCAU*	P=0	52.54	ICAM

Example 70

Incorporation of Cholesterol and Aminohexyl Modifications into 2'F Chimeric Oligonucleotides

The parent oligonucleotide, complementary to messenger 5 RNA of PKCa, is a chimeric phosphorothicate oligonucleotide having a central deoxyribonucleotide region that is "flanked" on both the 5' and the 3' sides with region that have 2'-deoxy-2'-fluoro nucleotides. These compound are alternately referenced as "gapped" oligonucleotides. The central "gapped" regions are shown in underlines in the Table below with the "flanks" (or alternatively the "wings") shown in unmodified type.

The oligonucleotides were synthesized, analyzed and purified as in Example 65, except for the following: 15 Concentration of all amidites was 0.1 M. Two 1- μ mol syntheses were performed for each oligonucleotide. The coupling step was extended 6 min, with periodic addition of fresh amidite; this extended coupling step was repeated once for the cholesterol amidite coupling. Trityl-on HPLC purification of 20 Compound ID #9532 was accomplished using a linear gradient of 5-40% AcCN over 50 min. After detritylation and purification by size-exclusion, it was analyzed using a similar analyticalscale HPLC procedure gel and capillary electrophoresis. Purified compound ID #9535 appears as two peaks by HPLC, Compound ID#s 9532 and 9534 contained 25 others as one peak. undeprotected 3'-hexylamine; probably about 20% according to previous results (phthalimide is the protecting group).

$$U_1 * = 3' - O - (CH_2)_6 - NH_2 U$$

 $U_2* = 3'-O-(CH_2)_6-NHCO-cholesterol U$

30 U_3 * = 2'-O-(CH₂)₆-NHCO-cholesterol U

Results are shown in the following Table were * indicated site of selected modification.

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	SEQUENCE	HPLC RETENTION TIME (min)
	UGUUCUC <u>GCTGGTGA</u> GUUUCAU	34.5
5	(SEQ ID NO:34) UGUUCUC <u>GCTGGTGA</u> GUUUCAU ₁ *	not determined
	UGUUCUCGCTGGTGAGUUUCAU2*	48 ·
	U3*GUUCUCGCTGGTGAGUUUCAU2*	46.5, 47.5
	U,*GUUCUCGCTGGTGAGUUUCAU	46.5, 47.5

EXAMPLE 71

10 5'-O-Dimethoxytrityl-3'-O-[hexyl-(omega-N-phtalimido)amino]uridine-2'-O-(2-cyanoethyl-N,N-diisopropyl)
Phosphoramidite

The 5'-dimethoxytrityl-3'-O-[hexyl-(omega-N-phthalimido)-amino]uridine (2 g, 2.6 mmol) was dissolved in 20 mL anhydrous 15 CH₂Cl₂. To this solution diisopropylaminotetrazolide (0.2 g, 1.16 mmol) and 2.0 mL 2-cyanoethyl-N,N,N',N'-tetraisopropyl phosphoramidite (6.3 mmol) were added and stirred overnight. TLC (1:1 EtOAc/hexane) showed complete disappearance of starting material. The reaction mixture was transferred with 20 CH₂Cl₂ and washed with saturated NaHCO₃ (100 mL), followed by saturated NaCl solution. The organic layer was dried over anhydrous Na₂SO₄ and evaporated to yield 3.8 g of a crude product, which was purified in a silica column (200 g) using 1:1 hexane/EtOAc to give 1.9 g (1.95 mmol, 74% yield) of the 25 desired phosphoramidite.

Sequence (3'→5')	Features of Sequence
GAU ₃ *CT	P=O, 2'-5' Linkage
GCG TGU3* CTG GG (SEQ ID NO:35)	P=O, 2'-5' Linkage
GCG UGU ₃ * CUG CG (SEQ ID NO:36)	P=O, 2'-5' Linkage, Chimera having 2'-OMe flanks

30 EXAMPLE 72

NMR and HPLC Analysis of 5'-3' versus 5'-2' Internucleotide Linkages and 2' substituents versus 3' substituents

The 400MHz 1 H spectrum of oligomer d(GAU₂*CT), where U₂= 2'-O-aminohexyluridine showed 8 signals between 7.5 and 9.0

ppm corresponding to the 8 aromatic protons. In addition, the anomeric proton of U* appears as a doublet at 5.9 ppm with $J_1'_{12}' = 7.5$ Hz, indicative of C2'-endo sugar puckering. corresponding 2'-5' linked isomer shows a similar structure 5 with $J_{1',2'} = 3.85$ Hz at 5.75 ppm, indicating an RNA type sugar puckering at the novel modification site favorable for hybridization to an mRNA target. The proton spectrum of the oligomer $d(GACU_3^*)$, where $U_3^*=3'-0$ - hexylamine, showed the expected 7 aromatic proton signals between 7.5 and 9.0 ppm and 10 the anomeric proton doublet at 5.9 ppm with $J_1', j' = 4.4$ Hz. This suggests more of a C3'-endo puckering for the 3'-Oalkylamino compounds compared to their 2' analogs. 31P NMR of these oligonucleotides showed the expected 4 and 3 signals from the internucleotide phosphate linkages for d(GAU*CT) and 15 d(GACU*), respectively. 3'-5' linked vs. 2'-5' linked have different retention times in RP HPLC and hence different lipophilicities, implying potentially different extent of interactions with cell membranes.

EXAMPLE 73

20 2'-5' Internucleotide Linked Oligonucleotide Conjugation Chemistry

The oligonucleotides 9274 and 9518 (20 ODs each) were dried and re-dissolved in 0.2M NaHCO3 buffer (200 µL) in an eppendorf tube. 5 mg of pyrene butyric acid-N-hydroxy 25 succinimide ester was dissolved in 400 µL anhydrous DMF. 200 µL of this solution was added to each of the eppendorf tubes and vortexed and left to stand overnight. The reaction mixtures were passed through a g-25 column to remove the excess reagent, an then purified by Reverse-Phase HPLC. The 30 retention times of the starting material and the conjugates are summarized below:

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Sequence (3'→5')	Retention Time (min)
GAU ₃ CT	23.85
GCG TGU3* CTG GG	24.99
pyrene conjugate of 9274	39.00
pyrene conjugate of 9518	40.05

HPLC conditions: Waters DeltaPak C-18 Delta Pak reverse-phase15m 300 Å column, 3.9x 300 mm size equipped with C-18 guard column; Solvent A: 50mM TEAA, pH 7.0.; Solvent B: acetonitrile; Gradient: 0-10 min., 5% acetonitrile; 10-60 5 min., 5% to 40% acetonitrile linear increase.

EXAMPLE 74

Tm Analysis of 2'-5' v. 3'-5' Connected Oligonucleotides and their Conjugates

Thermal melts were done as per the method of Procedure 10 B(1)(b): Oligonucleotide identity is as follows:

Oligonucleotide 31-A is a normal 3'-5' linked phosphodiester oligodeoxyribonucleotide of the sequence d(GGC TGU* CTG CG) (SEQ ID NO:37) where the * indicates the attachment site of either a 2'-aminolinker or a 2'-aminolinker 15 plus pyrene.

Oligonucleotide 31-B is a normal 3'-5' linked phosphodiester oligoribonucleotide of the sequence d(GGC TGU* CTG CG) where the * indicates the attachment site of either a 2'-aminolinker or a 2'-aminolinker plus pyrene. Each of the 20 ribonucleotides of the oligonucleotide, except the one bearing the * substituent, are 2'-O-methyl ribonucleotides.

Oligonucleotide 31-C is has 2'-5' linkage at the *
position in addition to either a 3'-aminolinker or a 3'aminolinker plus pyrene at this site. The remainder of the
25 oligonucleotide is a phosphodiester oligodeoxyribonucleotide
of the sequence d(GGC TGU* CTG CG). The base oligonucleotide
(no 2'-aminolinker) was not included in the study.

PCT/US94/10131

	OLIGONUCLEOTIDE	MODIFICATI	ON DN TARGET	'A	ΔTm TARGE	RNA T	AIM
5	31-A	none 2'-aminolinker plus pyrene	52.2 50.9 55.3	4.4	54.1° 55.5 55.5	O	
	31-B	none 2'-aminolinker plus pyrene	51.5 49.8 50.3	0.4	72.3 69.3 67.7	-1.6	
10	31-C	none 3'-aminolinker plus pyrene	NA 42.7 43.1	3	51.7 46.3	-5.4	

The 2'-5' linkages demonstrated a higher melting temperature against an RNA target compared to a DNA target.

EXAMPLE 75

15 2'-0-(Propylphthalimide) adenosine and 3'-0-(Propylphthal-imide) adenosine

To a solution of adenosine (20.0 g, 75 mmol) in dry dimethylformamide (550 ml) at 5°C was added sodium hydride (60% oil, 4.5 g, 112 mmol). After one hour, N-(3-20 bromopropyl)phthalimide (23.6 g, 86 mmol) was added and the temperature was raised to 30°C and held for 16 hours. Ice is added and the solution evaporated in vacuo to a gum. The gum was partitioned between water and ethyl acetate (4 x 300 ml). The organic phase was separated, dried, and evaporated in 25 vacuo and the resultant gum chromatographed on silica gel (95/5 CH₂Cl₂/MeOH) to give a white solid (5.7 g) of the 2'-O-(propylphthalimide)adenosine. Those fractions containing the 3'-O-(propylphthalimide)adenosine were rechromatographed on silica gel using the same solvent system.

Crystallization of the 2'-O-(propylphthalimide) adenosine fractions from methanol gave a crystalline solid, m.p. 123-124°C. 1 H NMR (400 MHz: DMSO-d₆) δ 1.70 (m, 2H, CH₂), 3.4-3.7 (m, 6H, C₅., CH₂, OCH₂, Phth CH₂), 3.95 (q, 1H, C₄.H), 4.30 (q, 1H, C₅.H), 4.46 (t, 1H, C₂.H), 5.15 (d, 1H, C₃.OH), 5.41 (t, 1H, 35 C₅.OH), 5.95 (d, 1H, C₁.H) 7.35 (s, 2H, NH₂), 7.8 (brs, 4H, Ar), 8.08 (s, 1H, C₂H) and 8.37 (s, 1H, C₈H). Anal. Calcd.

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 $C_{21}H_{22}N_6O_6$: C, 55.03; H, 4.88; N, 18.49. Found: C, 55.38; H, 4.85; N, 18.46.

Crystallization of the 3'-O-(propylphthalimide) adenosine fractions from $\rm H_2O$ afforded an analytical sample, m.p. 178-5 179°C. ¹H NMR (400 MHz: DMSO-d₆) δ 5.86 (d, 1H, H-1'),

EXAMPLE 76

2'-O-(Propylphthalimide)-No-benzoyladenosine

2'-O-(Propylphthalimide) adenosine was treated with benzoyl chloride in a manner similar to the procedure of 10 Gaffney, et al., Tetrahedron Lett. 1982, 23, 2257. Chromatography on silica gel (ethyl acetate-methanol) gives the title compound. Anal. Calcd. C₂₈H₂₆N₆O₇: C, 60.21; H, 4.69; N, 15.05. Found: C, 59.94; H, 4.66; N, 14.76.

EXAMPLE 77

15 2'-0-(Propylphthalimide)-5'-0-dimethoxytrityl-N-benzoyladenosine

To a solution of 2'-O-(propylphthalimide)-N⁶-benzoyl-adenosine (4.0 g) in pyridine (250 ml) was added 4,4'-dimethoxytrityl chloride (3.3 g). The reaction was stirred 20 for 16 hours. The reaction was added to ice/water/ethyl acetate, the organic layer was separated, dried, and concentrated in vacuo and the resultant gum chromatographed on silica gel (ethyl acetate-methanol triethylamine) to give the title compound. Anal. Calcd. C₄₉H₄₄N₆O₉: C, 68.36; H, 5.15; 25 N, 9.76. Found: C, 68.16; H, 5.03; N, 9.43.

EXAMPLE 78

N_6 -Benzoyl-5'-O-dimethoxytrityl-2'-(propylphthalimide)-3'-O-(N,N-diisopropyl- β -cyanoethyl)adenosine Phosphoramidite

2'-O-(Propylphthalimide)-5'-O-dimethoxytrityl-N⁶-30 benzoyladenosine was treated with (β-cyanoethoxy)chloro-N,N-diisopropyl)aminophosphane in a manner similar to the procedure of Seela, et al., Biochemistry 1987, 26, 2233. Chromatography on silica gel (EtOAc/hexane) gave the title compound as a white foam. ³¹P NMR (CD₃CN) 150.88, 151.22.

2'-0-(Propylamino) adenosine

A solution of 2'-O-(propylphthalimide) adenosine (8.8 g, 19 mmol), 95% ethanol (400 ml) and hydrazine (10 ml, 32 mmol) 5 was stirred for 16 hrs at room temperature. The reaction mixture was filtered and filtrate concentrated *in vacuo*. Water (150 ml) was added and acidified with acetic acid to pH 5.0. The aqueous solution was extracted with EtOAc (2 x 30 ml) and the aqueous phase was concentrated *in vacuo* to afford 10 7.1 g (95%) of the titled compound as a HOAc salt. ¹H NMR (200 MHz, DMSO-d₆) δ 1.70 (m, 2H, CH₂), 2.76 (m, 2H, CH₂NH₂), 3.55-3.67 (m, 4H, C₅,CH₂, OCH₂), 3.00 (q, 1H, C₄,H), 4.30 (q, 1H, C₃,H), 4.47 (t, 1H, C₂,H), 6.0 (d, 1H, C₁,H), 7.39 (s, 2H, NH₂) and 8.16 (s, 1H, C₂H).

15 EXAMPLE 80

2'-O-[3-(N-trifluoroacetamido)propyl]adenosine

A solution of 2'-O-(propylamino) adenosine in methanol (50 ml) and triethylamine (15 ml, 108 mmol) was treated with ethyl trifluoroacetate (18 ml, 151 mmol). The reaction was stirred 20 for 16 hrs and then concentrated in vacuo and the resultant gum chromatographed on silica gel (9/1, EtOAc/MeOH) to give the title 3.3 g of the compound (54%). m.p. 200-201°C (CH₃CN). ¹H NMR (200 MHz, DMSO-d₆) δ 1.7 (m, 2H, CH₂), 3.2 (m, 2H, CH₂NHCO), 3.37-3.67 (m, 4H, C₅.CH₂, OCH₂), 4.00 (q, 1H, C₃.H), 4.35 (q, 1H, C₃.H), 4.49 (t, 1H, C₂.H), 5.26 (d, 1H, C₃.OH), 5.43 (t, 1H, C₅.OH), 6.02 (d, 1H, C₁.H), 7.38 (s, 2H, NH₂), 8.16 (s, 1H, C₂H), 8.40 (s, 1H, C₈H), 9.34. (s, 1H, NHCO). ¹⁹F NMR (200 MHz, DMSO) δ -108.76. Anal. Calcd. for C₁₅H₁₉F₃N₆O₅: C, 42.86; H, 4.56; N, 19.99. Found: C, 43.10; H, 30 4.52; N, 20.03.

EXAMPLE 81

N⁶-(Dibenzoyl)-2'-0-[3-(N-trifluoroacetamido)propyl]adenosine

2'-0-[3-(N-trifluoroacetamido)propyl]adenosine is treated as per Example 76 using a Jones modification wherein 35 tetrabutylammonium fluoride is utilized in place of ammonia hydroxide in the work up. The crude product was purified using silica gel chromatography (EtOAc \rightarrow EtOAc/MeOH 1/1) to give the title compound. ¹H NMR (200 MHz, DMSO-d₆) δ 1.7 (m, 2H, CH₂), 3.2 (m, 2H, CH₂NHCO), 3.45-3.68 (m, 4H, C₅,CH₂, OCH₂), 5 4.03 (q, 1H, C₄,H), 4.37 (q, 1H, C₃,H), 4.53 (t, 1H, C₂,H), 5.18 (d, 1H, C₃,OH), 5.34 (d, 1H, C₃,OH), 6.17 (d, 1H, C₁,H), 7.45-7.83 (m, 1H, Ar), 8.73 (s, 1H, C₂H), 8.90 (s, 1H, C₈H), 9.37. (s, 1H, NHCO). ¹⁹F NMR (200 MHz, DMSO) δ -108.76. Anal. Calcd. for C₂₈H₂₇F₃N₆O₇: C, 55.41; H, 4.33; N, 13.37. Found: C, 10 55.16 H, 4.24; N, 13.09.

EXAMPLE 82

N⁶- (Dibenzoyl) - 5'-O- (dimethoxytrityl) - 2'-O-[3-(N-trifluoroacetamido)propyl] adenosine

4,4'-Dimethoxytrityl chloride (3.6 g, 10.0 mmol.) was added to a solution of N⁶-(dibenzoyl)-2'-O-[3-(N-trifluoro-acetamido) propyl) adenosine in pyridine (100 ml) at room temperature and stirred for 16 hrs. The solution was concentrated in vacuo and chromatographed on silica gel (EtOAc/TEA 99/1) to give the title compound. ¹H NMR (200 MHz, 20 DMSO-d₆) δ 1.74 (m, 2H, CH₂), 3.23 (m, 2H, CH₂NHCO), 3.60-3.73 (m, 10H, ArOCH₃, C₅.CH₂, OCH₂), 4.09 (m, 1H, C₄.H), 4.51 (q, 1H, C₃.H), 4.66 (t, 1H, C₂.H), 5.34 (d, 1H, C₃.OH), 6.17 (d, 1H, C₁.H), 6.80-7.83 (m, 23H, Ar), 8.64 (s, 1H, C₂H), 8.78 (s, 1H, C₈H), 9.35. (s, 1H, NHCO). ¹⁹F NMR (200 MHz, DMSO) δ -108.76. 25 Anal. Calcd. for C₄₉H₄₅F₃N₆O₉: C, 68.36; H, 5.15; N, 9,76. Found: C, 68.16 H, 5.03; N, 9.43.

EXAMPLE 83

 N^6 -(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-30 trifluoroacetamido)propyl]-3'-0-(N,N-diisopropyl- β -cyanoethyl)adenosine Phosphoramidite

A solution of N⁶-(dibenzoyl)-5'-O-(dimethoxytrityl)-2'-O[3-(N-trifluoroacetamido)propyl]adenosine in dry CH₂Cl₂ was
treated with bis-N,N-diisopropylamino cyanoethyl phosphite
35 (1.1 eqiv) and N,N-diisopropylaminotetrazolide (catalytic
amount) at room temperature for 16 hrs. The reaction was

concentrated in vacuo and chromatographed on silica gel (EtOAc/hexane/TEA 6/4/1) to give 0.5 g of the title compound.

31P NMR (200 MHz, CD₃CN) 150.87; (200 MHz, CDCL₃) 150.55, 150.83.

19F NMR (200 MHz, CD₃CN) 107.14, (200 MHz, CDCl₃) 5 102.67.

EXAMPLE 84

2'-0- (Butylphthalimide) adenosine

The title compound is prepared as per example 75, using N-(4-bromobutyl)phthalimide in place of the 1-bromopropane.

10 Chromatography on silica gel (EtOAC-MeOH) gives the title compound as a white solid. M.P. 199-200°C. Anal. Calcd. C₂₂H₂₄N₆O₆: C, 56.42; H, 5.16; N, 17.94. Found: C, 56.31; H, 5.04; N, 17.95.

The 3' compound can be separated from the mother liquors 15 as per Example 75 above. ^{1}H NMR (200 MHz, DMSO-d₆) δ 5.88 (d, 1H, C₁,H).

EXAMPLE 85

2'-O-(Butylphthalimide-No-benzoyladenosine)

Benzoylation of 2'-O-(butylphthalimide)adenosine as per 20 Example 76 gave the title compound. 1H NMR (200 MHz, DMSO- d_6) δ 6.18 (d, 1H, C_1 .H). Anal. Calcd. $C_{29}H_2N_6O_7$: C, 60.83; H, 4.93; N, 14.68. Found: C, 60.48; N, 14.41.

EXAMPLE 86

 $2'-0-(Butylphthalimide)-5'-0-dimethoxytrityl-N^6-25$ benzoyladenosine

The title compound was prepared from 2'-O-(butyl-phthalimide)-N⁶-benzoyladenosine as per Example 77. ¹H NMR (200 MHz, DMSO-d₆) δ 6.20 (d, 1H, C₁.H). Anal. Calcd. for C₅₀H₄₆N₆O₉: C, 68.64; H, 5.29; N, 9.60. Found: C, 68.47; H, 30 5.12; N, 9.37.

 N^6 -Benzoyl-5'-0-(dimethoxytrityl)-2'-0-(butylphthalimide)-3'-0-(N,N-diisopropyl- β -cyanoethyl)adenosine Phosphoramidite

The title compound was prepared from 2'-O-(butylphthal-5 imide)-5'-O-dimethoxytrityl-N⁶-benzoyladenosine as per Example 78. ³¹P NMR (CD₃CN) S 150.88, 151.22.

EXAMPLE 88

2'-O-(Pentylphthalimido)adenosine

The title compound was prepared as per Example 75, using 10 N-(5-bromopentyl)phthalimide. The crude material from the extraction was chromatographed on silica gel using CHCl₃/MeOH (95/5) to give a mixture of the 2' and 3' isomers. The 2' isomer was recrystallized from EtOH/MeOH 8/2. The mother liquor was rechromatographed on silica gel to afford the 3' 15 isomer.

2'-O- (Pentylphthalimido) adenosine: M.P. 159-160°C. Anal. Calcd. for $C_{23}H_{24}N_6O_5$: C, 57.26; H, 5.43; N, 17.42. Found: C, 57.03; H, 5.46; N, 17.33.

3'-O-(Pentylphthalimido)adenosine: 1 H NMR (DMSO- d_6) δ 5.87 20 (d, 1H, H-1').

EXAMPLE 89

2'-O- (Pentylphthalimido)-No-benzoyladenosine

Benzoylation of 2'-O-(pentylphthalimido)adenosine was achieved as per the procedure of Example 76 to give the title 25 compound. 1H NMR (DMSO-d₆): 6.10 (d, 1H, C₁, H). Anal. Calcd. for $C_{30}H_{30}N_6O_7$: C, 61.43; H, 5.16; N, 14.33. Found: C, 61.51; H, 4.97; N, 14.10.

EXAMPLE 90

2'-O-(Pentylphthalimido)-5'-O-(dimethoxytrityl)-N⁶-30 benzoyladenosine

The title compound was prepared from 2'-O-(pentyl-phthalimide)-N⁶-benzoyladenosine as per the procedure of Example 77. Chromatography on silica gel (ethylacetate, hexane, triethylamine), gave the title compound. Anal. Calcd.

for $C_{51}H_{48}N_6O_9$: C, 68.91; H, 5.44; N, 9.45. Found: C, 68.65; H, 5.45; N, 9.61.

EXAMPLE 91

 N^6 -Benzoyl-5'-0-(dimethoxytrityl)-2'-0-(pentylphthalimido)-3'-5 (N,N-diisopropyl- β -cyanoethyl) Phosphoramidite

The title compound was prepared from 2'-O-(pentyl-phthalimide)-5'-O-(dimethoxytrityl)-N⁶-benzoyladenosine as per the procedure of Example 78 to give the compound as a white foam.

10 EXAMPLE 92

2-Amino-2'-O-(propylphthalimide) adenosine and 2-Amino-3'-O-(propylphthalimide) adenosine

To a solution of 2-aminoadenosine (75 g, 281 mmol) in dry dimethylformamide (2000 ml) at room temperature was added 15 sodium hydride 60% (13.0 g, 325 mmol). After one hour N-(3-bromopropane)phthalimide (86.6 g, 3.23 mmol) was added and the temperature was raised to 40°C and held for 16 hrs. A further aliquot of sodium hydride 60% (6.2 g, 155 mmol) and N-(3-bromopropyl)phthalimide (40 g, 150 mmol) was added and heating 20 was continued for an additional 16 hrs. The reaction mixture was cooled, quenched with MeOH and concentrated in vacuo. The resultant gum was chromatographed on silica gel (CHCl₃/MeOH 9/1) to give 53.3 g of the 2' isomer (42%). Those fractions containing the 3' isomer were crystallized from EtOAc/MeOH to 25 yield 20 g of the 3' isomer (16%).

2-Amino-2'-O-(propylphthalimide) adenosine: ^{1}H NMR (200 MHz, DMSO-d₆) δ 1.82 (m, 2H, CH₂), 3.35-3.63 (m, 4H, OCH₂, C₅, H₂), 3.91 (m, 1H, C₄, H), 4.26 (m, 1H, C₃, H), 4.39 (m, 1H, C₂, H), 5.10 (d, 1H, C₃, OH), 5.44 (t, 1H, C₅, OH), 5.77 (s, 2H, 30 NH₂), 5.81 (d, 1H, C₁, H), 6.79 (bs, 2H, NH₂), 7.85 (bs, 4H, Ar) and 7.97 (s, 1H, C₈H). Anal. Calcd. for C₂₁H₂₃N₇O₆: C, 53.72;, H, 4.94; N, 20.88. Found: C, 53.70; H, 4.48; N, 20.56.

2-Amino-3'-O-(propylphthalimide) adenosine: m.p. 222-224°C, 1 H NMR (DMSO-d₆) δ 5.69 (d, 1H, H-1').

2'-0-(Propylphthalimide)guanosine

2-Amino-2'-O-(propylphthalimide)adenosine (38.6 g, 82 mmol) was dissolved in dimethylsulfoxide (DMSO, 1000 ml). 5 Sodium phosphate (0.1 M, 63 ml) and Tris buffer (0.5 M, 1200 ml) were added with stirring. The solution was adjusted to pH 7.0 with phosphoric acid. Adenosine deaminase (Sigma, 1.7 g) was added and the temperature raised to 35°C. The solution was stirred for 4 days with an additional aliquot of adenosine 10 deaminase (0.2 g) added every 24 hrs. The solution was maintained at pH 7.0 with the addition of phosphoric acid. The reaction was cooled, filtered and concentrated in vacuo. The resultant gum was crystallized from MeOH to give 31.6 of the title compound (82%). ¹H NMR (200 MHz, DMSO- d_6) δ 1.82 15 (m, 2H, CH₂), 3.45 (m, 4H, OCH₂, C₅.H₂), 3.85 (m, 1H, C₄.H), 4.26 (m, 2H, C₃.H, C₂.H), 5.09 (M, 2H, C₃.OH, C₅.OH), 5.80 (d, 1H, C_1,H), 6.47 (s, 2H, NH_2), 7.84 (s, 4H, Ar), 7.97 (s, 1H, C_8H) AND 10.65 (s, 1H, ArNH). Anal. Calcd. for $C_{21}H_{21}N_{6}O_{7}$: C, 53.72;, H, 4.94; N, 20.88. Found: C, 54.02; H, 4.73; N, 20 20.53.

EXAMPLE 94

2'-0-(Propylamino) guanosine

2'-O- (Propylphthalimide) guanosine was treated as per the procedure of Example 79 in water/methanol as the solvent to 25 give the title compound in 91% yield. ¹H NMR (200 MHz, DMSO-d₆) δ 1.80 (m, 2H, CH₂), 3.35-3.70 (m, 4H, OCH₂, C₅, H₂), 3.85 (m, 1H, C₄, H), 4.27 (m, 2H, C₃, H, C₂, H), 5.10 (m, 2H, C₃, OH, C₅, OH), 5.78 (d, 1H, C₁, H), 6.49 (s, 2H, NH₂) and 7.97 (s, 1H, C₈H).

EXAMPLE 95

30 2'-0-[3-(N-trifluoroacetamido)propyl]guanosine

2'-O-(Propylamino) guanosine was treated as per the procedure of Example 80 to give the title compound.

N²-(Dibenzoyl)-2'-0-[3-(N-trifluoroacetamido) propyl] guanosine
2'-0-[3-(N-trifluoroacetamido) propyl] guanosine was
treated as per the procedure of Example 81 to give the title
5 compound.

EXAMPLE 97

N²-(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-trifluoroacetamido)propyl]guanosine

N²-(Dibenzoyl)-2'-0-[3-(N-trifluoroacetamido)propyl]10 guanosine was treated as per the procedure of Example 82 to give the title compound.

EXAMPLE 98

 N^2 -(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-15 trifluoroacetamido)propyl]-3'-0-(N,N-diisopropyl- β -cyanoethyl)guanosine Phosphoramidite

N²-(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-trifluoroacetamido)propyl]guanosine was treated as per the procedure of Example 83 to give the title compound. ³¹P NMR 20 (CDCl₃) 150.49, 150.91. ¹⁹F NMR (CDCl₃) 102.71, 102.75.

EXAMPLE 99

2'-0-(Propylphthalimide)uridine and 3'-0-(Propylphthalimide)uridine

A solution of uridine-tin complex (prepared as per 25 Example 3) (48.2 g, 115 mmol) in dry DMF (150 ml) and N-(3-bromopropyl)phthalimide (46 g, 172 mmol) was heated at 130°C for 6 hrs. The crude product was chromatographed directly on silica gel CHCl₃/MeOH 95/5. The isomer ration of the purified mixture was 2'/3' 81/19. The 2' isomer was recovered by 30 crystallization from MeOH. The filtrate was rechromatographed on silica gel using CHCl₃→CHCl₃/MeOH (95/5) gave the 3' isomer as a foam.

2'-O- (Propylphthalimide) uridine: Analytical sample recrystallized from MeOH, m.p. 165.5-166.5°C, ¹H NMR (200 MHz, 35 DMSO-d₆) δ 1.87 (m, 2H, CH₂), 3.49-3.65 (m, 4H, C₂.H), 3.80-

3.90 (m, 2H, C₃,H₁C₄,H), 4.09 (m, 1H, C₂,H), 5.07 (d, 1h, C₃,OH), 5.16 (m, 1H, C₅,OH), 5.64 (d, 1H, CH=), 7.84 (d, 1H, C₁,H), 7.92 (bs, 4H, Ar), 7.95 (d, 1H, CH=) and 11.33 (s, 1H, ArNH). Anal. C₂₀H₂₁N₃H₈, Calcd. C, 55,69; H, 4.91;, N, 9.74. Found, C, 55.75; H, 5.12; N, 10.01.

3'-O-(Propylphthalimide)uridine: 1 H NMR (DMSO-d₆) δ 5.74 (d, 1H, H-1').

EXAMPLE 100

2'-O-(Propylamino)uridine

The title compound was prepared as per the procedure of Example 79 (90% yield). ^{1}H NMR (200 MHz, DMSO-d₆) δ 1.80 (m, 2H, CH₂), 3.57-3.66 (m, 4H, OCH₂,C₅,H₂), 3.88 (m, 2H, C₃,H,C₄,H), 4.15 (m, 1H, C₂,H), 5.64 (d, 1H, CH=), 5.82 (d, 1H, C₁,H), 7.02 (bs, 2H, NH₂) and 8.00 (d, 1H, C₈H).

15 **EXAMPLE 101**

2'-O-[3-(N-trifluoroacetamido)propyl]uridine

2'-O- (Propylamino) uridine was treated as per the procedure of Example 80 to give the title compound. ¹H NMR (200 MHz, DMSO-d₆) δ 1.75 (m, 2H, CH₂), 3.22-3.35 (m, 2H, 20 CH₂NH), 3.38-3.67 (m, 4H, OCH₂,C₅.H₂), 5.16 (m, 2H, C₃.OH,C₅.OH), 5.67 (d, 1H, CH=), 7.95 (d, 1H, CH=), 9.34 (bs, 1H, NHCO) and 11.36 (s, 1H, ArNH). ¹⁹F NMR (200 MHz, DMSO) 108.76. Anal. Calcd. for C₁₄H₁₈F₃N₃O₇: C, 42.31; H, 4.56; N, 10.60. Found: 42.56; H, 4.61; N, 10.25.

25 EXAMPLE 102

5'-O-(dimethoxytrity1)-2'-O-[3-(N-trifluoroacetamido)propy1]-uridine

2'-O-[3-(N-trifluoroacetamido)] propyl] uridine was treated as per the procedure of Example 82 to give the title compound. 30 ^{1}H NMR (200 MHz, DMSO-d₆) δ 1.95 (m, 2H, CH₂), 3.42-3.80 (m, 12H, 2HOCH, OCH, CH), 4.03 (m, 2H, CH), 4.45 (m, 1H, COH)

12H, $ArOCH_3$, OCH_2 , C_5 , H_2), 4.03 (m, 2H, C_2 .H), 4.45 (m, 1H, C_3 .OH), 5.30 (d, 1H, CH=), 5.91 (s, 1H, C_1 .H), 6.83-7.43 (m, 13H, ArH), 7.90 (bs, 1H, NHCO), 8.10 (d, 1H, CH=) and 10.3 (s, 1H, ArNH). ¹⁹F NMR (200 MHz, DMSO) 102.75.

5'-O- (Dimethoxytrityl) -2'-O- [3-(N-trifluoroacetamido) propyl] -3'-O- (N,N-diisopropyl- β -cyanoethyl) uridine Phosphoramidite

5'-O-(Dimethoxytrityl)-2'-O-[3-(N-trifluoroacetamido)-propyl]uridine was treated as per the procedure of Example 83 to give the title compound. ³¹P NMR (200 MHz, CD₃CN) 154.87, 155.62. ¹⁹F NMR (200 MHz, CD₃CN) 107.19.

EXAMPLE 104'

10 2'-0-(Propylphthalimide)cytidine and 3'-0-(Propylphthalimide)cytidine

The title compounds were prepared as per the procedure of Example 75.

2'-O- (propylphthalimide) cytidine: ¹H NMR (200 MHz, DMSO-15 d₆) δ 5.82 (d, 1H, C₁.H).

3'-O- (propylphthalimide) cytidine: ¹H NMR (200 MHz, DMSO-d₆) δ 5.72 (d, 1H, C₁.H).

EXAMPLE 105

2'-0-(Propylamino)cytidine

20 2'-O-(Propylphthalimide) cytidine was treated as per the procedure of Example 79 to give the title compound. ^{1}H NMR (200 MHz, DMSO-d₆) δ 5.82 (d, 1H, C₁.H).

EXAMPLE 106

2'-0-[3-(N-trifluoroacetamido)propyl]cytidine

2'-0-(Propylamino) cytidine was treated as per the procedure of Example 80 to give the title compound. ¹⁹F NMR (DMSO-d₆) 108.73.

EXAMPLE 107

N'- (Benzoyl) -2'-0-[3-(N-trifluoroacetamido)propyl] cytidine

30 2'-O-[3-(N-trifluoroacetamido) propyl] cytidine was treated as per the procedure of Example 81 to give the title compound.

19F NMR (DMSO-d₆) 108.79.

N⁴-(Benzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-trifluoroacet-amido)propyl]cytidine

N⁴-(Benzoyl)-2'-0-[3-(N-trifluoroacetamido)propyl]-5 cytidine was treated as per the procedure of Example 82 to give the title compound.

EXAMPLE 109

N⁴-(Benzoyl)-5'-0-(dimethoxytrityl)-2'-0-[3-(N-trifluoroacet10 amido)propyl]-3'-0-(N,N-diisopropyl-β-cyanoethyl)cytidine
Phosphoramidite N⁴-(Benzoyl)-5'-0-(dimethoxytrityl)-2'-0[3-(N-trifluoroacetamido)propyl]cytidine was treated as per
the procedure of Example 83 to give the title compound. ³¹P
NMR (CDCl₃) 150.03, 151.36. ¹⁹F NMR (CDCl₃) 102.76.

15 EXAMPLE 110

2'-O-[3-(Imidizo-1-yl)propyl]adenosine

The title 2'-O-substituted compound was prepared as per the procedure of Example 75 using 1-(4-bromopropyl)imidazole. 1 H NMR (200 MHz, DMSO-d₆) δ 6.02 (d, 1H, C₁.H).

20 The corresponding 3'-O- substituted compound can be separated from the mother liquors as per Example 75. ¹H NMR (200 MHz, DMSO-d₆) δ 5.91 (d, 1H, C₁·H).

EXAMPLE 111

2'-0-[4-(Imidizo-1-yl)butyl)] adenosine and 3'-0-[4-(Imidizo-1-25 yl)butyl] adenosine

A solution of adenosine (6.97 g, 26 mmol) in dry DMF (250 ml) was treated with NaH 60% (3.9 g, 97 mmol) for 15 minutes. N-(4-Bromobutyl)imidazoline acetate salt (9.4 g, 2.1 equiv. HOAc, 29 mmol) was added and the reaction mixture stirred at 30 room temperature for 2 days. The reaction was quenched (MeOH) and concentrated in vacuo to a semi-solid. The solid was dissolved in water (250 ml) and extracted with EtOAc (2 x 100 ml). The aqueous phase was concentrated in vacuo and chromatographed on silica gel using EtOAc/MeOH (6/4) to give 35 a mixture of 2' and 3' isomers (70/30, 62%). Crystallization

of this solid from EtOAc/MeOH 8/2 gave the 2' isomer. Concentration of the mother liquors and subsequent crystallization from MeOH afforded the 3' isomer in 12% yield.

2'-O-[4-(Imidizo-1-yl)] butyl] adenosine: ¹H NMR (200 MHz, 5 DMSO-d₆) δ 6.01 (d, 1H, C₁,H).

3'-O-[4-(Imidizo-1-yl)] butyl]adenosine: m.p. 183-184°C. ¹H NMR (DMSO-d₆) δ 5.89 (d, 1H, H-1').

EXAMPLE 112

2'-0-[4-(Imidizo-1-yl)butyl]-N6-benzoyladenosine

The title compound was prepared 2'-0-[4-(imidizo-1-yl)-butyl]adenosine as per Example 76.

EXAMPLE 113

2'-0-[4-(Imidizo-1-yl)butyl]-5'-0-dimethoxytrityl-N6-benzoyl-adenosine

The title compound was prepared from N⁶-benzoyl-2'-O-[4-(imidizo-1-yl)butyl]adenosine as per Example 77.

EXAMPLE 114

 N^6 -Benzoyl-5'-0-(dimethoxytrityl)-2'-0-[4-(imidizo-1-yl)-(butyl]-3'-0-(N,N-diisopropyl- β -cyanoethyl)adenosine Phosphor-20 amidite

The title compound was prepared from 2'-O-[4-(imidizo-1-yl)] butyl]-5'-O-dimethoxytrityl-N⁶-benzoyladenosine as per Example 78. ³¹P NMR (CDCl₃) 150.48, 151.06.

EXAMPLE 115

25 2-Amino-2'-0-[4-(imidizo-1-yl)butyl] adenosine and 2-Amino-3'0-[4-(imidizo-1-yl)butyl] adenosine

The title compounds are prepared as per Example 111 with the substitution of 2-aminoadenosine for adenosine. The isomer ratio of 2'/3' was 74/26.

2-Amino-2'-O-[4-(imidizo-1-yl)butyl] guanosine: ¹H NMR (200 MHz, DMSO- d_6) δ 5.84 (d, 1H, C_1 -H).

2-Amino-3'-O-[4-(imidizo-1-yl)butyl] guanosine: ¹H NMR (200 MHz, DMSO-d₆) δ 5.68 (d, 1H, C₁·H).

2'-0-[4-(Imidazo-1-yl)butyl]guanosine

In the manner described in Example 93, 2-amino-2'-O-[4-(imidazo-1-yl)butyl]adenosine in sodium phosphate (0.1 M) and 5 Tris buffer (0.5 M) at pH 7.0 is deaminated with adenosine deaminase (Sigma) at 35°C to give the title compound.

EXAMPLE 117

N2-(Dibenzoyl)-2'-0-[4-(imidazo-1-yl)butyl]guanosine

2'-0-[4-(Imidazo-1-yl)butyl]guanosine was treated as per 10 the procedure of Example 81 to give the title compound.

EXAMPLE 118

N²-(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[4-(imidazo-1-yl)butyl]guanosine

N²-(Dibenzoyl)-2'-0-[4-(imidizo-1-yl)butyl]guanosine was 15 treated as per the procedure of Example 82 to give the title compound.

EXAMPLE 119

 N^2 -(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[4-(imidazo-1-20 yl)butyl]-3'-0-(N,N-diisopropyl- β -cyanoethyl)guanosine Phosphoramidite

 N^2 -(Dibenzoyl)-5'-0-(dimethoxytrityl)-2'-0-[4-(imidazo-1-yl)butyl]guanosine was treated as per the procedure of Example 83 to give the title compound. ^{31}P NMR (CDCl₃) 150.67, 150.80

25 EXAMPLE 120

2'-0-[4-(Imidizo-1-yl)butyl]cytidine and 3'-0-[4-(imidizo-1-yl)-butyl]cytidine

The title compounds were prepared as per the procedure of Example 111 with the substitution of cytidine for adenosine.

- 30 The crude product was chromatographed on silica gel using EtOAc/MeOH/HOAc. The isomer ratio of 2'/3' was 60/40.
 - 2'-O-[4-(Imidizo-1-yl)] butyl] cytidine: ¹H NMR (200 MHz, DMSO-d_s) δ 5.88 (d, 1H, C₁.H).

3'-O-[4-(Imidizo-1-yl)] butyl] cytidine: ¹H NMR (DMSO-d₆) δ 5.78 (d, 1H, H'1')

EXAMPLE 121

2'-0-[4-(Imidizo-1-yl)butyl]-N'-benzoylcytidine

The title compound was prepared 2'-0-[4-(imidizo-1-yl)-butyl]cytidine as per Example 76.

EXAMPLE 122

2'-0-[4-(Imidizo-1-yl)butyl]-5'-0-dimethoxytrityl-N'-benzoyl-cytidine

The title compound was prepared from N⁴-benzoyl-2'-O-[4-(imidizo-1-yl)butyl]cytidine as per Example 77.

EXAMPLE 123

N'-Benzoyl-5'-0-(dimethoxytrityl)-2'-0-[4-(imidizo-1-yl)-(butyl]-3'-0-(N,N-diisopropyl- β -cyanoethyl)cytidinePhosphor15 amidite

The title compound was prepared from 2'-O-[4-(imidizo-1-yl)butyl]-5'-O-dimethoxytrityl-N4-benzoylcytidine as per Example 78. 31P NMR (CDCl₃) 150.39, 150.92.

EXAMPLE 124

20 3-Benzyloxymethyl-5-methyluridine

A solution of 5-methyluridine (51.6, 200 mmol) in DMF (100 ml) was treated with sodium hydride (60% oil, 9.9 g, 248 mmol). After 1 hr at room temperature, benzylchloromethyl ether (34.4 g, 220 mmol) was added dropwise and the reaction 25 mixture heated to 35°C for 4 hrs. The reaction mixture was cooled, quenched with MeOH (5 ml), filtered and concentrated in vacuo. The resultant gum was partitioned between EtOAc/sat. NaCl, separated, dried and the organic phase concentrated whereupon the title compound crystallized. ¹H 30 NMR (DMSO-d₆) δ 5.83 (d, 1H, C₁,-H).

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EXAMPLE 125

3-Benzyloxymethyl-5-methyl-2'-0-(4-chlorobutyl)uridine

A solution of 3-benzyloxymethyl-5-methyluridine (25.9 g, 68 mmol) was treated with NaH (60% oil, 7.9 g, 200 mmol) for 5 1 hr at room temp. 1-Bromo-4-chlorobutane (19.7 g, 115 mmol) was added slowly and the reaction stirred at room temp. for 16 hrs. The reaction was then quenched (MeOH), concentrated in vacuo and the residue chromatographed on silica gel CHCl₃/MeOH 95/5 to give the title compound in 41% yield. ¹H 10 NMR (DMSO-d₆) δ 5.87 (d, 1H, C₁.H).

EXAMPLE 126

3-Benzyloxymethyl-5-methyl-2'-0-[4-(imidazo-1-yl)butyl]uridine

To a solution of imidazole (6.8 g, 100 mmol) in DMF (150 ml) was added NaH (60% in oil, 4 g, 100 mmol). After 4 hrs, 15 this suspension was added to a suspension of 3-benzyloxymethyl-5-methyl-2'-0-(4-chlorobutyl)uridine (8.7 g, 18 mmol), NaI (2.6 g) and DMF (200 ml). The reaction was stirred 16 hrs at room temp., quenched with MeOH, concentrated in vacuo and the residue chromatographed on 20 silica gel CHCl₃/MeOH 85/15→50/50 to give the title compound in 31% yield. ¹H NMR (DMSO-d₆) δ 5.87 (d, 1H, C₁.H).

EXAMPLE 127

2'-0-[4-(Imidazo-1-yl)butyl]-5-methyluridine

A solution of 3-benzyloxymethyl-5-methyl-2'-O-[4-25 (imidazo-1-yl)butyl]uridine (5.4 g, 10 mmol) in MeOH (200 ml) was hydrogenated with 20% palladium hydroxide (9.0 g) at 1 atmosphere H₂ for 16 hrs. The reaction was filtered through celite and the mother liquor concentrated to afford the product in 83% yield. ¹H NMR (DMSO-d₆) δ 5.86 (d, 1H, C₁.H).

5'-0-(Dimethoxytrityl)-2'-0-[4-(imidazo-1-yl)butyl]-5-methyluridine

5-Methyl-2'-O-[4-(imidazo-1-yl)butyl]uridine was treated 5 as per the procedure of Example 77 to give the title compound in 91% yield. 1 H NMR (DMSO- d_6) δ 5.86 (d, 1H, C_1 H).

EXAMPLE 129

5'-0- (Dimethoxytrity1) -2'-0- [4-(imidazo-1-yl)buty1] -3'-0- (N,N-diisopropy1- β -cyanoethy1) -5-methyluridine Phosphoramidite

5'-O-(Dimethoxytrityl)-5-methyl-2'-O-[4-(imidazo-1-yl)butyl]uridine was treated as per the procedure of Example 78 to give the title compound in 83% yield. ¹H NMR (DMSO-d₆) δ 5.85 (d, 1H, C₁.H). ³¹P NMR (DMSO-d₆) 153.99, 154.45.

Example 130

15 Effect Of Derivatization On Melting Temperature

Melting temperatures (Tm) for a variety of modified oligonucleotides according to the invention were determined according the method set forth in Procedure B(1)(b). The resulting data is shown in Tables 5-8, wherein all listed 20 oligonucleotides contain 2'-deoxy sugars and phosphodiester backbones unless otherwise indicated.

Table 5 shows changes in Tm that result by modifying the 2'-position of adenosine nucleotides to include a variety of substituent groups. For example, the first entry shows that 25 the introduction of 2'-O-aminopropyl adenosine substituents at two adenosine nucleotides resulted in a heteroduplex having a Tm that is 1.13°C greater than the heteroduplex formed by an unmodified oligonucleotide having the same sequence.

Table 6 shows changes in Tm that result by modifying one 30 or more 2'- and 3'-positions of the indicated parent oligonucleotides to include linkers and pyrene conjugates. These modifications were made to terminal and internal (i.e., non-terminal) positions within the oligonucleotide. The notation "PS" indicates the presence of a phosphorothicate 35 backbone, and the notation "2'-5' linkage" indicates 2'-5'

intersugar linkages. As can be seen from the first entry, the introduction of a 2'-aminopentoxy linker to a single, internal adenosine nucleotide of the parent oligonucleotide resulted in a Tm decrease of 3.1°C whereas the introduction of a 2'-5 pyrenyl-aminopentoxy conjugate resulted in a Tm increase of 3.3°C.

Tables 7a-c show changes in Tm that result by modifying one or more positions of the indicated parent oligonucleotides to include linkers and cholate or cholesterol conjugates. As 10 can be seen from the first entry, the introduction of an aminolink linker at the 3'-position of a single terminal nucleotide of the parent phosphorothicate oligonucleotide resulted in a Tm increase of 1.3°C and the addition of a cholate group to the linker resulted in a further Tm increase 15 of 1.1°C.

Table 8 shows changes in Tm that result by modifying one or more 3'-positions of the indicated parent oligonucleotides to include linkers and PEG conjugates. The entries under the heading "PEG vs Unmodified" represent preliminary findings.

20 As can be seen, the first entry indicates that introduction of an aminolink linker at the 3'-position of a single terminal nucleotide of the parent phosphodiester oligonucleotide resulted in a Tm increase of 0.9°C and that the addition of a PEG 550 group to the linker did not change the Tm of the resulting product.

TABLE 5 - THE INFLUENCE OF THE 2'-0-dA MODIFICATIONS ON THE HETERODUPLEX STABILITY

TARGETS			Tm/Modification	lication		
	2'0-propyl -dA	2'-O-amino propyl-dA	2'-0-butyl -dA	2'-0-pentyl -dA	2'-0-amino pentyl-dA	2'-O-butyl- imidazole-dA
GAGCTCCCAGGC (SEQ ID NO:38) (2 mod.)		+1.13	99'0-	56'0-	+0.60	-0.1
CGACTATGCAAGTAC (SEQ ID NO:39) (1 mod.)			-1.01	-1.50	00.36	+0.14
CGACTATGCAAAAAC (SEQ ID NO:40) (3 mod.)		+0.98	+0.51	+0.07	+0.85	•
ACCGAGGATCATGTCGTA CGC (SEQ ID NO:41) (5 mod.)	-0.36	-0.40	-0.56	-0.92	-1.11	-0.45
Average	-0.2 +-0.7	-0.15+-1.15	-0.4+-1.0		-0.9+-0.6 -0.35+-1.1	-0.31+-0.6

	TABELE (TABELE 6 - EFFECT OF PYRENE AND LINKERS ON TM VS DNA	ND LINKE	RS ON TM	VS DNA					
					Pyrene vs Linker Only	ine S Only	Linker Only vs Unmodified	Only s diffed	Pyrene vs Unmodifi	Pyrene vs Unmodified
Pyrene Conjugate	Linker Only	Unmodified Parent	# Mods	Pos. of Mods.	dTm	dTm/ Mod.	dTm	dTm/ Mod.	фTm	dTm/ Mod.
2'-pyrenyl-amino- pentoxy-dA	2'-amino-pentoxy dA	wild type DNA	1	internal	+6.5	+6.5	-3.1	-3.1	+3.3	+3.3
2'-pyrenyl-amino- hexoxy-dU	2'-amino-hexoxy dU	wild type DNA	-	internal	+4.4	+4.4	-1.2	-1.2	+3.1	+3.1
2'-pyrenyl-amino-hexoxy dU in 2'-0-methyl backggrounde	2'-amino-hexoxy dU in _2'-0-methyl background	fully 2'-0-methyl with	-:	internal	+0.5	+0.5	-1.7	1-7	-1.2	-1.2
2'-pyrenyl-amino- pentoxy-dA	2'-amino-pentoxy dA	wild type DNA	-	internal	+4.5	+4.5	-2.4	-2.4	+2.0	+2.0
2'-pyrenyl-amino- pentoxy-dA	2'-amino-pentoxy dA	wild type DNA	2	internal	+8.3	+4.2	-3.7	-1.8	+4.6	+2.3
3'-pyrenyl-amino- hexoxy-dU	3'-amino-hexoxy dU	wild type DNA	-	3'- term.	+2.8	+2.8	-0.7	-0.7	+2.2	+2.2
3'-amino-linked pyrene (PS)	3'-aminolink (PS)	fully PS DNA	-	3'- term.	+0.7	+0.7	+0.6	+0.6	+1.2	+1.2
3'-pyrenyl-amino- hexoxy-dU {2'-5' linkage)	3'-amino-hexoxy-dU (2'- 5' linkage)	DNA with dU replacing 3'-amino- hexoxy-dU	-	internal	+0.4	+0.4	-8.9	-8.9	-8.5	-8.5

	TABLE 7a - EFFECT	TABLE 78 - EFFECT OF CHOLATE OR CHOLESTEROL AND LINKERS ON TH VS RNA	STEROL ,	AND LINKE	RS ON T	W VS RNA				
					Cholate/ cholesterol vs Linker Only	ate/ terol Only	Linker Only vs Unmodified	Only s diffied	Cholate/ cholesterol vs Unmodified	ate/ sterol s dified
Cholate Conjugate	Linker Only	Unmodified Parent	# Mods	Pos. of Mods.	đТm	dTm/ Mod.	đТm	dTm/ Mod.	dTm	dTm/ Mod.
3'amino-linked cholate (PS)	3'-aminolink (PS)	fully PS DNA	1	3'- term.	+1.1	+1.1	+1.3	+1.3	+2.5	+2.5
3'-amino-linked cholesterol (PS)	3'-aminolink (PS)	fully PS DNA	1	3′- term.	7		+1.3	+1.3	7	
3'-amino-linked cholate	3'-amino-link	wild type DNA	-	3'- term.	+2.0	+2.0	-0.3	-0.3	+1.7	+1.7
5'-amino-linked cholate	5'-aminolink	wild type DNA	-	5'- term.	+0.4	+0.4	+0.2	+0.2	+0.6	+0.6
5'-amino-linked cholate	5'-aminolink	wild type DNA	-	5′- term.	:		:		+0.1	+0.1
5'-amino-linked cholate (PS)	5'-aminolink (PS)	fully PS DNA	1	5'- term.	:		t		+0.9	+0.9
2'cholesterol-amino- hexoxy dU¹	•	DNA with dU in position 1	-	5-term.	•		1		-1.3	
2'cholesterol-amino- hexoxy dU¹	•	DNA with dU in position 10		internal	ŧ		;		-4.5	
2'cholesterol-amino- hexoxy dU (PS) ¹		fully PS DNA with dU in position 1	1	5′. term.	1		:		-3.7	
2'-adamantane-amino- hexoxy dU (PS)¹	•	fully PS DNA with dU in position 1	-	5'- term.					0.2	

	TABLE 7b - EFFECT OF (TABLE 7b - EFFECT OF CHOLATE OR CHOLESTEROL AND LINKERS ON TM VS RNA (CONT.)	TOL AND	LINKERS	ON TH VS	RNA (CC	NT.)			
					Cholate/ cholesterol vs Linker Only	ate/ terol s Only	Linker Only vs Unmodiffed	Only s diffed	Cholate/ cholesterol vs Unmodified	ate/ iterol s dified
Cholate Conjugate	Linker Only	Unmodified Parent	# Mods	Pos. of Mods.	ďTm	dTm/ Mod.	đđm	dTm/ Mod.	dTm	dTm/ Mod.
2'-eiconsenic-amino- hexoxy dU (PS)¹		fully PS DNA with dU in position 1		5′- term.	1		ì		-1.3	
3'-cholesterol-amino- hexoxy dU (PS)¹	ŧ	fully PS DNA with dU in position 20	-	3'- term.	:		1		-0.5	
2'cholesterol-amino- hexoxy dU-in position 1 3'-cholesterol-amino- hexoxy dU position 20 (PS) ¹		fully PS DNA with dU in positions 1 and 20	2	both ends	1		1	!	-5.7	
2'-cholesterol-amino- hexoxy dU in position 1 3'-PEG-amino-hexoxy dU position 20 (PS)	-	fully PS DNA with dU in positions 1 and 20	7	both ends	ł		ŀ		-5.2	-2.6
3'-cholesterol-amino- hexoxy dU1	•	DNA with dU in position 20	-	3'. term.	:		1		+0.6	
2'-cholesterol-amino- hexoxy dU in position 1 3'-cholesterol-amino- hexoxy dU position 20'	1	DNA with dU in positions 1 and 20	7	both ends	!		!		-3.9 -3.9	

¹ Synthesis via cholesterol amidite.

	TABLE 7c - EFFECT	TABLE 7c - EFFECT OF CHOLATE OR CHOLESTEROL AND LINKERS ON THE VS DNA	STEROL /	AND LINKE	RS ON T	NS DN	1			
					Cholate/ cholesterol vs Linker Only	ste/ terol Only	Linker Only vs Unmodified	er Only vs nodified	Cholate/ cholesterol vs Unmodified	ate/ iterol s diffed
Cholate Conjugate	Linker Only	Unmodified Parent	# Mods	Pos. of Mods.	dTm	dTm/ Mod.	dTm	dTm/ Mod.	ďľm	dTm/ Mod.
2'-cholate-amino- pentoxy dA	2'-amino-pentoxy dA	wild type DNA	1	internal	-2.6	-2.6	-2.4	-2.4	-5.1	-5.1
2'-cholate-amino- pentoxy dA	2'-amino-pentoxy dA	wild type DNA	2	internal	-2.5	-1.2	-3.7	-1.8	-6.2	-3.1
3'-amino-linked cholate (PS)	3'-aminolink (PS)	fully PS DNA	1	3'- term.	+0.4	+0.4	+0.6	+0.6	+1.0	+1.0
3'-amino-linked cholesterol (PS)	3'-aminolink (PS)	fully PS DNA	1	3′- term.			+0.6	+0.6		
3'-amino-linked cholate	3'-amino-link	wild type DNA	1	3'- term.	+1.4	+1.4	-0.5	-0.5	+0.9	+ 0.9
5'-amino-linked cholate	5'-aminolink	wild type DNA		5′- term.	+0.9	+0.9	+0.1	+0.1	+0.9	+ 0.9
5'-amino-linked cholate	5'-aminolink	wild type DNA	-	5'- term.	ŀ		1		+0.9	6.0+
5'-amino-linked cholate (PS)	5'-aminolink (PS)	fully PS DNA	1	5′- term.	:				+1.0	+1.1

	AT.	TABLE 8 - EFFECT OF PEG AND LINKERS ON THE VS RNA	AND LINK	ERS ON 1	W VS RNA					
					PEC vs Linker	PEG vs Linker only	Linker only vs Unmodified	Linker only vs Unmodified	PE v Unmo	PEG vs Unmodified
PEG Conjugate	Linker only	Unmodified Parent	# Mods	Pos. of Mods.	dTm	dTm/ Mod.	dTm	dTm/ Mod.	dTm	dTm/ Mod.
3'-aminolinked PEG 550	3'-aminolink	wild type DNA		3'- term.	+0.0	+ 0.0	+0.9	+0.9	+0.9	+0.9
3'-aminolinked PEG 2000	3'-aminolink	wild type DNA	-	3'- term.	-0.4	-0.4	+0.9	+0.9	+0.5	+0.5
3'-aminolinked PEG 5000	3'-aminolink	wild type DNA	. . .	- 3'. term.	-0.3	-0.3	÷0.9	+0.9	+0.6	+0.6
3'-aminolinked PEG 550 (PS)	3'-aminolink (PS)	fully PS DNA	1	3'- term.	+0.2	+0.2	+1.4 +0.3	+1.4 +0.3	+1.0	+1.0
3'aminolinked PEG 2000 (PS)	3'-aminolink (PS)	fully PS DNA	1	3'. term.	+0.6	+0.6	+1.4 +0.3	+1.4 +1.4	+1.4	+1.4
3'-aminolinked PEG 5000 (PS)	3'-aminolink (PS)	fully PS DNA	1	3′- term.	-0.0	-0.0	+1.4 +0.3	+1.4 +0.3	+0.8	+0.8
2'-cholesterol-amino- hexoxy dU in position 1 3'-PEG-amino-hexoxy dU position 20 (PS) ¹	:	fully PS DNA with dU in positions 1 and 20	2	both ends	1		t		-5.2	-2.6

¹ Synthesis via cholesterol amidite.

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SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANTS: ISIS Pharmaceuticals, Inc., et al.
 - (ii) TITLE OF INVENTION: AMINE-DERIVATIZED NUCLEOSIDES AND OLIGONUCLEOSIDES
 - (iii) NUMBER OF SEQUENCES: 41
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz and Norris
 - (B) STREET: One Liberty Place 46th Floor
 - (C) CITY: Philadelphia
 - (D) STATE: PA
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 19103
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: WordPerfect 5.1
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: N/A
 - (B) FILING DATE: Herewith
 - (C) CLASSIFICATION:
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/117,363
 - (B) FILING DATE: 9/3/94
 - (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 07/854,634
 - (B) FILING DATE: 7/1/92
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Lucci, Joseph
 - (B) REGISTRATION NUMBER: 33,307
 - (C) REFERENCE/DOCKET NUMBER: ISIS-1691
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 215-568-3100
 - (B) TELEFAX: 215-568-3439
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 base pairs
 - (B) TYPE: nucleic acid

- 101 -

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
	GGCTGACTGC G	11
(2)	INFORMATION FOR SEQ ID NO:2:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
	CTGTCTCCAT CCTCTTCACT	20
(2)	INFORMATION FOR SEQ ID NO:3:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
	TTGCTTCCAT CTTCCTCGTC	20
(2)	INFORMATION FOR SEQ ID NO:4:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
	TGGGAGCCAT AGCGAGGC	18
(2)	INFORMATION FOR SEQ ID NO:5:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(vi) SPOURNCE DESCRIPTION: SEO ID NO:5:	

PCT/US94/10131

	CTGTCTCCAT CCTCTTCACT	20
(2)	INFORMATION FOR SEQ ID NO:6:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 16 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
	TCCAGGTGTC CGCATC	16
(2)	INFORMATION FOR SEQ ID NO:7:	•
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
	GGACCGGAAG GTACGAG	17
(2)	INFORMATION FOR SEQ ID NO:8:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
	CCTGGCCTTC CATGCTC	17
(2)	INFORMATION FOR SEQ ID NO:9:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
	CCUGGCCUUC CAUGCUC	17
(2)	INFORMATION FOR SEQ ID NO:10:	

	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
	CCCA	AGGCUCA GA	12
(2)	INFO	RMATION FOR SEQ ID NO:11:	
`	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	
	GAGC	CUCCCAG GC	12
(2)	INFO	RMATION FOR SEQ ID NO:12:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:12:	
	CAUG	GCUGCAG CC	12
(2)	INFO	RMATION FOR SEQ ID NO:13:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:13:	
	GCCI	TTTCGCG ACCCAACACU	20
(2)	INFO	RMATION FOR SEQ ID NO:14:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	

	(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	•
	GCGTGUCTGC G	10
(2)	INFORMATION FOR SEQ ID NO:15:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:	
	UGGGAGCCAT AGCGAGGC	18
(2)	INFORMATION FOR SEQ ID NO:16:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 18 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:	
	TGGGAGCCAU AGCGAGGC	18
(2)	INFORMATION FOR SEQ ID NO:17:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:	
	UCTGAGTAGC AGAGGAGCTC	20
(2)	INFORMATION FOR SEQ ID NO:18:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	
	TICCCONNECT CCCNTCCCTC N	21

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(2)	INFORMATION FOR SEQ ID NO:19:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:	
	UGCGTTTGCT CTTCTTCT CG	22
(2)	INFORMATION FOR SEQ ID NO:20:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
•	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:	
	UGCATCCCCC AGGCCACCAT	20
(2)	INFORMATION FOR SEQ ID NO:21:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:	
٠	UCCCGCCTGT GACATGCATT	20
(2)	INFORMATION FOR SEQ ID NO:22:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:	
	GUTCTCGCTG GTGAGTTTCA	20
(2)	INFORMATION FOR SEQ ID NO:23:	
	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 19 base pairs	

	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:	
	UUGGGAGCCA TAGCGAGGC	19
(2)	INFORMATION FOR SEQ ID NO:24:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:	
	CCAAGCCUCA GA	12
(2)	INFORMATION FOR SEQ ID NO:25:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:	
	CCAGGCUCAG AT	12
(2)	INFORMATION FOR SEQ ID NO:26:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:	
	UGCATCCCCC AGGCCACCA	19
(2)	INFORMATION FOR SEQ ID NO:27:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:	
	UCAGTGCCTG CGCCGCGCTC G	21
(2)	INFORMATION FOR SEQ ID NO:28:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
٠	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:	
	UCCGTCATCG CTCCTCAGGG	20
(2)	INFORMATION FOR SEQ ID NO:29:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:	
	UTTAGGATTC GTGCTCATGG	20
(2)	INFORMATION FOR SEQ ID NO:30:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
	UGTTCTCGCT GGTGAGTTTC A	21
(2)	INFORMATION FOR SEQ ID NO:31:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
	TGCATCCCCC AGGCCACCAII	20

(2)	INFORMATION FOR SEQ ID NO:32:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
	GTGCTCATGG TGCACGGTCU	20
(2)	INFORMATION FOR SEQ ID NO:33:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	٠
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
	UGCATCCCCC AGGCCACCAU	20
(2)	INFORMATION FOR SEQ ID NO:34:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 22 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
	UGUUCUCGCT GGTGAGUUUC AU	22
(2)	INFORMATION FOR SEQ ID NO:35:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:	
	GCGTGUCTGG G	11
(2)	INFORMATION FOR SEQ ID NO:36:	
	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs	

		(B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:36:	
	GCGU	JGUCUGC G	11
(2)	INFO	RMATION FOR SEQ ID NO:37:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:37:	
	GGCT	rguctgc g	11
(2)	INFO	RMATION FOR SEQ ID NO:38:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:38:	
	GAGC	CTCCCAG GC	12
(2)	INFO	RMATION FOR SEQ ID NO:39:	
		SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:39:	
	CGAC	CTATGCA AGTAC	15
(2)	INFO	RMATION FOR SEQ ID NO:40:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 15 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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		•
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
	CGACTATGCA AAAAC	15
(2)	INFORMATION FOR SEQ ID NO:41:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:	
	ACCGAGGATCATGTCGTACGC	21

WHAT IS CLAIMED IS:

1. A compound comprising a plurality of linked nucleosides, wherein:

each nucleoside includes a ribofuranosyl sugar portion and a base portion; and

at least one of said nucleosides bears at a 2'-0-position, a 3'-0-position, or a 5'-0-position a substituent having formula:

$$-R_A-N(R_{1a})(R_{1b})$$

where:

 R_A is alkyl having from 1 to about 10 carbon atoms or $(CH_2-CH_2-Q)_x$;

 R_{1a} and R_{1b} , independently, are H, R_2 , or an amine protecting group, have formula $C(X)-R_2$, $C(X)-R_A-R_2$, $C(X)-Q-R_A-R_2$, or, together, complete a cycloalkyl or cycloaryl moiety having two to six carbon atoms and one or two nitrogen atoms; and

 R_2 is a steroid molecule, a reporter molecule, a lipophilic molecule, a reporter enzyme, a peptide, a protein, or has formula $-Q-(CH_2CH_2-Q-)_x-R_3$;

X is 0 or S;

each Q is, independently, is NH, O, or S;

x is 1 to about 200;

 R_3 is H, R_A , C(O)OH, C(O)OR, C(O)R, RA-N3, or R_A -NH₂; R_4 is Cl, Br, I, SO₂R₅ or has structure:

m is 2 to 7; and

R₅ alkyl having 1 to about 10 carbon atoms.

- 2. The compound of claim 1 wherein more than one of said nucleosides bear said substituent at a 2'-O-position, a 3'-O-position, or a 5'-O-position.
- 3. The compound of claim 1 wherein $R_{\rm A}$ is $(CH_2)_{\rm n}$ where n is an integer from 1 to about 10.

- 4. The compound of claim 3 wherein n is 6.
- 5. The compound of claim 1 wherein said R_{1a} and R_{1b} , together, are phthalimido.
- 6. The compound of claim 1 wherein R_{1a} is H and R_{1b} is $C(O)-(CH_2)_n-R_2$ where n is an integer from 1 to about 10.
- 7. The compound of claim 1 wherein R_{1a} is H and R_{1b} is R_{2} .
- 8. The compound of claim 1 wherein R_{1a} is H and R_{1b} is $C(0) O R_2$.
- 9. The compound of claim 1 wherein R_{1a} and R_{1b} both are alkyl.
- 10. The compound of claim 1 wherein R_{1a} is H and R_{1b} is $C(O) (CH_2)_n R_2$ where n is an integer from 1 to about 10.
- 11. The compound of claim 10 wherein R_2 has formula $(CH_2)_5$ -NH-.
- 12. The compound of claim 1 wherein R_{1a} is H and R_{1b} is $C(S)-NH-R_2$.
- 13. The compound of claim 1 wherein R2 includes pyrene, fluorescein, dinitrophenyl, cholesterol, acridine.
- 14. The compound of claim 1 wherein R_{1a} is H and R_{1b} is $C\left(O\right)$ R_{2} .
- 15. The compound of claim 14 wherein R_2 has formula $-0-(CH_2CH_2-0-)_x-R_3$.

- 16. The compound of claim 1 wherein a 3'-position of at least one of said nucleosides is linked to a 5'-position of an adjacent nucleoside.
- 17. The compound of claim 1 wherein a 2'-position of at least one of said nucleosides is linked to a 5'-position of an adjacent nucleoside.
- 18. A compound comprising a plurality of linked nucleosides, wherein:

each nucleoside includes a ribofuranosyl sugar portion and a base portion; and

at least one of said nucleosides bears at a 2'-O-position, a 3'-O-position, or a 5'-O-position a substituent having formula:

$$-R_{A}-N(R_{1a})(R_{1b})$$

where:

 R_{λ} is alkyl having from 1 to about 7 carbon atoms; and

 R_{1a} and R_{1b} are H.

- 19. The compound of claim 18 wherein R_A is propyl.
- 20. A compound comprising a plurality of linked nucleosides, wherein:

each nucleoside includes a ribofuranosyl sugar portion and a base portion; and

at least one of said nucleosides bears at a 2'-0-position, a 3'-0-position, or a 5'-0-position a substituent having formula:

$$-R_A-N(R_{1a})(R_{1b})$$

where:

 $R_{\rm A}$ is alkyl having from 1 to about 7 carbon atoms; and

 $R_{\rm la}$ and $R_{\rm lb},$ together, complete a cycloaryl moiety having two to six carbon atoms and one or two nitrogen atoms.

- 21. The compound of claim 20 wherein $R_{\rm A}$ is butyl and $R_{\rm 1a}$ and $R_{\rm 1b}$, together, are imidazole.
- 22. A nucleoside comprising a ribofuranosyl sugar portion and a base portion, wherein said nucleoside bears at a 2'-O-position, a 3'-O-position, or a 5'-O-position a substituent having formula:

$$-R_A-N(R_{1a})(R_{1b})$$

where:

 R_{λ} is alkyl having from 1 to about 10 carbon atoms:

 R_{1a} and R_{1b} , independently, are H, R_2 , or an amine protecting group or have formula $C(X)-R_2$, $C(X)-R_A-R_2$, $C(X)-Q-R_A-R_2$; and

 R_2 is a steroid molecule, a reporter molecule, a lipophilic molecule, a reporter enzyme, a peptide, a protein, or has formula $-Q-(CH_2CH_2-Q-)_x-R_3$;

X is O or S;

each Q is, independently, is NH, O, or S;

x is 1 to about 200;

 $\rm R_3$ is H, $\rm R_A$, C(O)OH, C(O)OR_A, C(O)R_4, R_A-N_3, or $\rm R_A-NH_2;$ and

 R_4 is Cl, Br, I, SO_2R_s or has structure:

m is 2 to 7; and

R, alkyl having 1 to about 10 carbon atoms.

- 23. The compound of claim 22 wherein R_A is $(CH_2)_n$ where n is an integer from 1 to about 10.
 - 24. The compound of claim 22 wherein n is 6.
- 25. The compound of claim 22 wherein said R_{1a} and R_{1b} , together, are phthalimido.

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- 26. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $C(O)-(CH_2)_n-R_2$ where n is an integer from 1 to about 10.
- 27. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $R_{2}\,.$
- 28. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $C(O) O R_2$.
- 29. The compound of claim 22 wherein R_{1a} and R_{1b} both are alkyl.
- 30. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $C(O) (CH_2)_n R_2$ where n is an integer from 1 to about 10.
- 31. The compound of claim 30 wherein R_2 has formula $-(CH_2)_5-NH-.$
- 32. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $C\left(S\right)-NH-R_{2}.$
- 33. The compound of claim 22 wherein R_2 includes pyrene, fluorescein, dinitrophenyl, cholesterol, acridine.
- 34. The compound of claim 22 wherein R_{1a} is H and R_{1b} is $C(O) R_2$.
- 35. The compound of claim 34 wherein R_2 has formula $-O-(CH_2CH_2-O-)_x-R_3$.
- 36. A method for modulating the production of a protein by an organism comprising contacting an organism with a compound of claim 1.
- 37. A method for modulating the production of a protein by an organism comprising contacting an organism with a compound of claim 22.

- 38. A method of treating an animal having a disease characterized by undesired production of protein comprising contacting said animal with a compound of claim 1.
- 39. A method of treating an animal having a disease characterized by undesired production of protein comprising contacting said animal with a compound of claim 22.
- 40. A method for detecting the presence or absence of an RNA in a biological sample suspected of containing said RNA comprising contacting said sample with a compound of claim 1.
- 41. A method for detecting the presence or absence of an RNA in a biological sample suspected of containing said RNA comprising contacting said sample with a compound of claim 22.

INTERNATIONAL SEARCH REPORT

Insurational application No. PCT/US94/10131

A. CLASSIFICATION OF SUBJECT MATTER IPC(6) :C07H 21/00, 21/02, 21/04; C12Q 1/68				
US CL: 435/6; 536/24.3, 25.3 According to International Patent Classification (IPC) or to both national classification and IPC				
B. FIELDS SEARCHED				
Minimum documentation searched (classification system followed by classification symbols)				
U.S. : 435/6; 536/24.3, 25.3				
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched				
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)				
APS, CAS				
C. DOCUMENTS CONSIDERED TO BE RELEVANT				
Category*	Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.	
Υ	US, A, 4,910,300 (URDEA ET A entire document.	AL.) 20 March 1990, see	1-41	
Y	SCIENCE, Vol. 238, issued 04 Dec "Generation of a Hybrid Sequence Deoxyribonuclease", pages 1401-1	e-Specific Single Stranded	1-41	
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Further documents are listed in the continuation of Box C. See patent family annex.				
Special extegories of cited documents: A* document defining the general state of the art which is not considered to be of particular relevance.		"T" later document published after the inte- date and not in conflict with the applica principle or theory underlying the inve	tion but cited to understand the	
	lier document published on or after the international filing date	"X" document of particular relevance; the considered novel or cannot be consider		
cite	cument which may throw doubts on priority claim(s) or which is and to establish the publication date of another citation or other scial reason (as specified)	when the document is taken alone "Y" document of particular relevance; the	claimed invention cannot be	
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	sument published prior to the international filing date but later than priority date claimed	eal filing date but later than '&' document member of the same patent family		
Date of the actual completion of the international search 07 DECEMBER 1994		Date of mailing of the international warch report 27 DEC 1994		
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT		Authorized officer SCOTT HOUTTEMAN Why 3c? for		
Washington, D.C. 20231 Facsimile No. (703) 305-3230		Telephone No. (703) 308-0196	$^{-}$	